

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214965_at	0.034221	Consensus includes gb:AF070574.1 /DEF=Homo sapiens clone 24819 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387945 /UG=Hs.22316 Homo sapiens clone 24819 mRNA sequence	AF070574		NP_689552
214972_at	0.0174	meningioma expressed antigen 5 (hyaluronidase)	AU144791	Hs.5734	
214988_s_at	0.016934	Consensus includes gb:X63071.1 /DEF=H:sapiens mRNA for novel DNA binding protein. /FEA=mRNA /GEN=DBP-5 /PROD=DBP-5 protein /DB_XREF=gi:30487 /UG=Hs.92909 SON DNA binding protein	X63071		NP_620305
214992_s_at	0.011725	Consensus includes gb:AD000092 /DEF=Homo sapiens DNA from chromosome 19p13.2 cosmid R31240, R30272 and R28549 containing the EKLF, GCDH, CRTC, and RAD23A genes, genomic sequence /FEA=CDS_1 /DB_XREF=gi:1905905 /UG=Hs.307251 Homo sapiens DNA from chromosome 19p13.2 cosmid R31240, R30272 and R28549 containing the EKLF, GCDH, CRTC, and RAD23A genes, genomic sequence	AD000092		
215000_s_at	0.001116	Consensus includes gb:AL117593.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G082 (from clone DKFZp564G082). /FEA=mRNA /DB_XREF=gi:5912157 /UG=Hs.103419 fasciculation and elongation protein zeta 2 (zygin II)	AL117593		NP_005093
215004_s_at	0.034317	Hypothetical human protein; Similar to novel proteins encoded by (AC004447) KIAA0365 [Homo sapiens] and (Y08265) DAN26 [Homo sapiens]; Homo sapiens chromosome 19, cosmid F23858, complete sequence.	AC004475		

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215030_at	0.046651	Consensus includes gb:AK023187.1 /DEF=Homo sapiens cDNA FLJ13125 fis, clone NT2RP3002877. /FEA=mRNA /DB_XREF=gi:10435001 /UG=Hs.287551 Homo sapiens cDNA FLJ13125 fis, clone NT2RP3002877	AK023187		
215031_x_at	0.013088	hypothetical protein FLJ20552	BG420893	Hs.69554	NP_060346
215032_at	0.004341	Consensus includes gb:AK022442.1 /DEF=Homo sapiens cDNA FLJ12380 fis, clone MAMMA1002556. /FEA=mRNA /DB_XREF=gi:10433838 /UG=Hs.300934 Homo sapiens cDNA FLJ12380 fis, clone MAMMA1002556	AK022442		
215038_s_at	7.58E-04	Consensus includes gb:AF049103.1 /DEF=Homo sapiens Huntingtin interacting protein mRNA, partial cds. /FEA=mRNA /PROD=Huntingtin interacting protein /DB_XREF=gi:3319281 /UG=Hs.306184 Huntingtin interacting protein B	AF049103		NP_054878
215043_s_at	0.022988	Consensus includes gb:X83301.1 /DEF=H.sapiens SMA5 mRNA. /FEA=mRNA /GEN=SMA5 /DB_XREF=gi:603029 /UG=Hs.324728 SMA5	X83301		
215047_at	1.58E-04	Consensus includes gb:AL080170.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C091 (from clone DKFZp434C091); partial cds. /FEA=mRNA /GEN=DKFZp434C091 /PROD=hypothetical protein /DB_XREF=gi:5262639 /UG=Hs.51692 DKFZP434C091 protein	AL080170		NP_056246

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215049_x_at	0.005852	Consensus includes gb:Z22969.1 /DEF=H.sapiens mRNA for M130 antigen cytoplasmic variant 1. /FEA=mRNA /PROD=M130 antigen cytoplasmic variant 1 /DB_XREF=gi:312143. /UG=Hs.74076 CD163 antigen	Z22969		NP_004235
215051_x_at	5.03E-04	allograft inflammatory factor 1	BF213829	Hs.76364	NP_116573
215063_x_at	0.001886	Consensus includes gb:AL390149.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547B026 (from clone DKFZp547B026). /FEA=mRNA /DB_XREF=gi:9368883 /UG=Hs.283862 Homo sapiens mRNA; cDNA DKFZp547B026 (from clone DKFZp547B026)	AL390149		
215087_at	7.58E-04	Consensus includes gb:AL109730.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 68600. /FEA=mRNA /DB_XREF=gi:5689835 /UG=Hs.306331 Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 68600	AL109730		
215088_s_at	1.26E-04	ESTs, Highly similar to C560_HUMAN Succinate dehydrogenase cytochrome B560 subunit, mitochondrial precursor (QPS1) (CII-3) (Succinate dehydrogenase complex subunit C) (Succinate-ubiquinone oxidoreductase cytochrome B large subunit) (CYBL) [H.sapiens]	BG110532	Hs.355964	
215118_s_at	0.040938	Homo sapiens translocation associated fusion protein IRTA1/IGA1 (IRTA1/IGHA1) mRNA, complete cds	AW519168	Hs.367852	
215127_s_at	2.05E-04	RNA binding motif, single stranded interacting protein 1	AL517946	Hs.241567	NP_058523

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215158_s_at	1.26E-04	Consensus includes gb:AK022531.1 /DEF=Homo sapiens cDNA FLJ12469 fis, clone NT2RM1000874, highly similar to Homo sapiens death effector domain-containing testicular molecule mRNA. /FEA=mRNA /DB_XREF=gi:10433972 /UG=Hs.169681 death effector domain-containing	AK022531		NP_127491
215175_at	3.26E-04	Consensus includes gb:AB023212.1 /DEF=Homo sapiens mRNA for KIAA0995 protein, partial cds. /FEA=mRNA /GEN=KIAA0995 /PROD=KIAA0995 protein /DB_XREF=gi:4589633 /UG=Hs.225967 KIAA0995 protein	AB023212		NP_055797
215176_x_at	0.022708	immunoglobulin kappa constant	AW404894	Hs.306357	
215193_x_at	0.004341	Consensus includes gb:AJ297586.1 /DEF=Homo sapiens mRNA for MHC class II antigen (HLA-DRB1 gene), DRB1*0402 allele. /FEA=CDS /GEN=HLA-DRB1 /PROD=MHC class II antigen /DB_XREF=gi:10185079 /UG=Hs.279930 major histocompatibility complex, class II, DR beta 3	AJ297586		NP_002115
215204_at	0.010205	ESTs, Weakly similar to 2109260A B cell growth factor [Homo sapiens] [H.sapiens]	AU147295	Hs.288575	
215210_s_at	0.003862	Consensus includes gb:S72422.1 /DEF=E2k=alpha-ketoglutarate dehydrogenase complex dihydrolipoyl succinyltransferase human, fetal brain, mRNA, 2987 nt. /FEA=mRNA /GEN=E2k /PROD=alpha-ketoglutarate dehydrogenase complex dihydrolipoyl succinyltransferase /DB_XREF=gi:632883 /UG=Hs.296348 E2k	S72422		

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215221_at	1.26E-04	Consensus includes gb:AK025064.1 /DEF=Homo sapiens cDNA: FLJ21411 fis, clone COL03986. /FEA=mRNA /DB_XREF=gi:10437503 /UG=Hs.306758 Homo sapiens cDNA: FLJ21411 fis, clone COL03986	AK025064		
215222_x_at	0.016934	Consensus includes gb:AK023406.1 /DEF=Homo sapiens cDNA FLJ13344 fis, clone OVARC1002050, highly similar to Homo sapiens mRNA for actin binding protein ABP620. /FEA=mRNA /DB_XREF=gi:10435330 /UG=Hs.321395 Homo sapiens cDNA FLJ13344 fis, clone OVARC1002050, highly similar to Homo sapiens mRNA for actin binding protein ABP620	AK023406		NP_149033
215223_s_at	0.00228	superoxide dismutase 2, mitochondrial	W46388	Hs.372783	NP_000627
215236_s_at	0.008185	ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase)	AV721177	Hs.114765	NP_665801
215273_s_at	0.001054	Consensus includes gb:AK024982.1 /DEF=Homo sapiens cDNA: FLJ21329 fis, clone COL02464. /FEA=mRNA /DB_XREF=gi:10437412 /UG=Hs.158196 transcriptional adaptor 3 (ADA3, yeast homolog) like (PCAF histone acetylase complex)	AK024982		NP_597815
215275_at	0.013223	EST375211 MAGE resequences, MAGH Homo sapiens cDNA, mRNA sequence.	AW963138		
215285_s_at	1.41E-04	putative homeodomain transcription factor 1	AA927671	Hs.123637	NP_006599
215287_at	0.003171	Homo sapiens ELISC-1 mRNA, partial cds	AA975427	Hs.128434	
215313_x_at	0.007779	major histocompatibility complex, class I, C	AA573862	Hs.77961	NP_002107

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215318_at	0.028552	Consensus includes gb:AL049782.1 /DEF=Novel human gene mapping to chromosome 13. /FEA=mRNA /DB_XREF=gi:4902604 /UG=Hs.184938 Novel human gene mapping to chromosome 13	AL049782		
215336_at	0.005039	Consensus includes gb:AK002166.1 /DEF=Homo sapiens cDNA FLJ11304 fis, clone PLACE1009997, weakly similar to Rattus norvegicus A-kinase anchoring protein AKAP 220 mRNA. /FEA=mRNA /DB_XREF=gi:7023877 /UG=Hs.232076 A kinase (PRKA) anchor protein 11	AK002166		NP_652761
215338_s_at	1.26E-04	natural killer-tumor recognition sequence	AI688640	Hs.241493	NP_005376
215343_at	0.034221	Consensus includes gb:AF070587.1 /DEF=Homo sapiens clone 24741 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387960 /UG=Hs.25770 Homo sapiens clone 24741 mRNA sequence	AF070587		
215359_x_at	7.58E-04	GIOT-2 for gonadotropin inducible transcription repressor-2	AI758888	Hs.251371	
215366_at	0.041055	Consensus includes gb:AL353943.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761E0611 (from clone DKFZp761E0611). /FEA=mRNA /GEN=DKFZp761E0611 /PROD=hypothetical protein /DB_XREF=gi:7669982 /UG=Hs.284191 Homo sapiens mRNA; cDNA DKFZp761E0611 (from clone DKFZp761E0611)	AL353943		
215373_x_at	0.005039	Consensus includes gb:AK022213.1 /DEF=Homo sapiens cDNA FLJ12151 fis, clone MAMMA1000431. /FEA=mRNA /DB_XREF=gi:10433560 /UG=Hs.287493 hypothetical protein FLJ12151	AK022213		

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215375_x_at	0.025245	Consensus includes gb:AK023938.1 /DEF=Homo sapiens cDNA FLJ13876 fis, clone THYRO1001401. /FEA=mRNA /DB_XREF=gi:10436033 /UG=Hs.287604 Homo sapiens cDNA FLJ13876 fis, clone THYRO1001401	AK023938		
215383_x_at	0.013223	Consensus includes gb:AL137312.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761K23121 (from clone DKFZp761K23121). /FEA=mRNA /DB_XREF=gi:6807797 /UG=Hs.293681 Homo sapiens mRNA; cDNA DKFZp761K23121 (from clone DKFZp761K23121)	AL137312		
215392_at	0.002418	AU148154 MAMMA1 Homo sapiens cDNA clone MAMMA1002744 3', mRNA sequence.	AU148154		
215399_s_at	1.26E-04	amplified in osteosarcoma	AI683900	Hs.76228	NP_006803
215404_x_at	0.016934	Consensus includes gb:AK024388.1 /DEF=Homo sapiens cDNA FLJ14326 fis, clone PLACE4000247. /FEA=mRNA /DB_XREF=gi:10436762 /UG=Hs.306678 Homo sapiens cDNA FLJ14326 fis, clone PLACE4000247	AK024388		
215424_s_at	5.03E-04	SKI-interacting protein	AV689564	Hs.79008	NP_036377
215434_x_at	0.0174	hypothetical protein DJ328E19.C1.1	AV684285	Hs.323463	NP_060410
215438_x_at	8.68E-04	G1 to S phase transition 1	BE906054	Hs.2707	NP_002085
215470_at	0.026891	Consensus includes gb:U21915.1 /DEF=Human chromosome 5q13.1 clone 5G8 mRNA. /FEA=mRNA /DB_XREF=gi:736412 /UG=Hs.14658 Human chromosome 5q13.1 clone 5G8 mRNA	U21915		

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215492_x_at	0.040938	Consensus includes gb:AL035587 /DEF=Human DNA sequence from clone RP3-475N16 on chromosome 6p12.3-21.2. Contains the genes for CTG4A, pre-T cell receptor alpha, a novel protein similar to RPL7A (60S ribosomal protein L7A) and the 3' end of gene KIAA0240. Contains ESTs, STSs, GSSs an... /FEA=mRNA_2 /DB_XREF=gi:6002306 /UG=Hs.169002 Human pre TCR alpha mRNA, partial cds	AL035587		
215498_s_at	2.05E-04	mitogen-activated protein kinase kinase 3	AA780381	Hs.180533	NP_659732
215499_at	3.26E-04	mitogen-activated protein kinase kinase 3	AA780381	Hs.180533	NP_659732
215511_at	0.003171	Consensus includes gb:U19345.1 /DEF=Homo sapiens AR1 (TCF20) mRNA, partial cds. /FEA=mRNA /GEN=TCF20 /PROD=AR1 /DB_XREF=gi:2924754 /UG=Hs.201668 transcription factor 20 (AR1)	U19345		
215524_x_at	0.013088	immunoglobulin heavy constant mu	AW966434	Hs.300697	
215548_s_at	0.003171	Consensus includes gb:AB020724.1 /DEF=Homo sapiens mRNA for KIAA0917 protein, partial cds. /FEA=mRNA /GEN=KIAA0917 /PROD=KIAA0917 protein /DB_XREF=gi:4240322 /UG=Hs.27023 vesicle transport-related protein	AB020724		NP_057247
215549_x_at	0.00423	Consensus includes gb:AC005587 /DEF=Homo sapiens PAC clone RP5-988G15 from 7q33-q35 /FEA=CDS_2 /DB_XREF=gi:4156166 /UG=Hs.272525 Homo sapiens PAC clone RP5-988G15 from 7q33-q35	AC005587		

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215566_x_at	0.013088	Consensus includes gb:AK024724.1 /DEF=Homo sapiens cDNA: FLJ21071 fis, clone CAS01789, highly similar to AF098668 Homo sapiens acyl-protein thioesterase mRNA. /FEA=mRNA /DB_XREF=gi:10437080 /UG=Hs.283655 lysophospholipase II	AK024724		NP_009191
215582_x_at	0.006603	Consensus includes gb:AK022303.1 /DEF=Homo sapiens cDNA FLJ12241 fis, clone MAMMA1001274. /FEA=mRNA /DB_XREF=gi:10433670 /UG=Hs.287503 Homo sapiens cDNA FLJ12241 fis, clone MAMMA1001274	AK022303		
215588_x_at	0.003171	Consensus includes gb:AK024958.1 /DEF=Homo sapiens cDNA: FLJ21305 fis, clone COL02124. /FEA=mRNA /DB_XREF=gi:10437382 /UG=Hs.287658 Homo sapiens cDNA: FLJ21305 fis, clone COL02124	AK024958		
215599_at	0.040938	Consensus includes gb:X83300.1 /DEF=H.sapiens SMA4 mRNA. /FEA=mRNA /GEN=SMA4 /DB_XREF=gi:603028 /UG=Hs.289103 SMA4	X83300		
215600_x_at	0.005852	Consensus includes gb:AK022174.1 /DEF=Homo sapiens cDNA FLJ12112 fis, clone MAMMA1000043. /FEA=mRNA /DB_XREF=gi:10433510 /UG=Hs.288793 Homo sapiens cDNA FLJ12112 fis, clone MAMMA1000043	AK022174		

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215602_at	6.68E-04	Consensus includes gb:AK024456.1 /DEF=Homo sapiens mRNA for FLJ00048 protein, partial cds. /FEA=mRNA /GEN=FLJ00048 /PROD=FLJ00048 protein /DB_XREF=gi:10440425 /UG=Hs.289034 Homo sapiens mRNA for FLJ00048 protein, partial cds	AK024456		
215603_x_at	8.94E-04	gamma-glutamyltransferase 1	AI344075	Hs.284380	
215604_x_at	0.013088	Consensus includes gb:AK023783.1 /DEF=Homo sapiens cDNA FLJ13721 fis, clone PLACE2000450. /FEA=mRNA /DB_XREF=gi:10435820. /UG=Hs.289035 Homo sapiens cDNA FLJ13721 fis, clone PLACE2000450	AK023783		
215608_at	0.033793	ESTs	AU144378	Hs.296650	
215623_x_at	0.005339	Consensus includes gb:AK002200.1 /DEF=Homo sapiens cDNA FLJ11338 fis, clone PLACE1010720, highly similar to Homo sapiens mRNA for chromosome-associated polypeptide-C. /FEA=mRNA /DB_XREF=gi:7023928 /UG=Hs.50758 SMC4 (structural maintenance of chromosomes 4, yeast)-like 1	AK002200		NP_005487
215628_x_at	0.001116	Consensus includes gb:AL049285.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564M193 (from clone DKFZp564M193). /FEA=mRNA /DB_XREF=gi:4500045 /UG=Hs.302053 Homo sapiens mRNA; cDNA DKFZp564M193 (from clone DKFZp564M193)	AL049285		
215629_s_at	0.013088	ESTs, Weakly similar to LEU2_HUMAN Leukemia associated protein 2 [H.sapiens]	AA905286	Hs.380846	
215633_x_at	0.041055	leukocyte specific transcript 1	AV713720	Hs.88411	
215646_s_at	3.26E-04	Homo sapiens versican Vint isoform, mRNA, partial cds	R94644	Hs.397468	
215667_x_at	0.016934	postmeiotic segregation increased 2-like 2	AI375694	Hs.177548	NP_777590

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215671_at	0.007779	AU144792 HEMBA1 Homo sapiens cDNA clone HEMBA1002973 3', mRNA sequence.	AU144792		
215684_s_at	5.07E-04	Consensus includes gb:AL096741.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586O0223 (from clone DKFZp586O0223). /FEA=mRNA /DB_XREF=gi:5419897 /UG=Hs.49005 hypothetical protein	AL096741		NP_115580
215694_at	0.018973	Consensus includes gb:AK022348.1 /DEF=Homo sapiens cDNA FLJ12286 fis, clone MAMMA1001768, weakly similar to CELL DIVISION CYCLE PROTEIN 48 HOMOLOG MJ1156. /FEA=mRNA /DB_XREF=gi:10433725 /UG=Hs.5555 hypothetical protein MGC5347	AK022348		NP_076968
215696_s_at	1.26E-04	Consensus includes gb:BC001404.1 /DEF=Homo sapiens, clone IMAGE:3139772, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3139772) /DB_XREF=gi:12655104 /UG=Hs.5716 KIAA0310 gene product	BC001404		
215698_at	3.26E-04	Consensus includes gb:AF007135.1 /DEF=Homo sapiens clone 23592 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852610 /UG=Hs.76272 retinoblastoma-binding protein 2	AF007135		
215716_s_at	0.033381	Consensus includes gb:L14561 /DEF=Homo sapiens plasma membrane calcium ATPase isoform 1 (ATP2B1) gene, alternative splice products, partial cds /FEA=mRNA_2 /DB_XREF=gi:4165324 /UG=Hs.78546 ATPase, Ca++ transporting, plasma membrane 1	L14561		

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215719_x_at	0.033381	Consensus includes gb:X83493.1 /DEF=H.sapiens mRNA for FasApo-1 (clone pCRTM11-FasdeltaTM). /FEA=mRNA /DB_XREF=gi:971457 /UG=Hs.82359 tumor necrosis factor receptor superfamily, member 6	X83493		NP_690616
215735_s_at	0.002418	Consensus includes gb:AC005600 /DEF=Homo sapiens chromosome 16, P1 clone 109-9G (LANL), complete sequence /FEA=CDS_1 /DB_XREF=gi:3522919 /UG=Hs.90303 tuberous sclerosis 2	AC005600		
215739_s_at	6.68E-04	Consensus includes gb:AJ003062.1 /DEF=Homo sapiens mRNA for protein encoded by Saccharomyces cerevisiae SPC98 homologue. /FEA=mRNA /DB_XREF=gi:3152381 /UG=Hs.9884 spindle pole body protein	AJ003062		NP_006313
215749_s_at	0.009199	Consensus includes gb:AK001574.1 /DEF=Homo sapiens cDNA FLJ10712 fis, clone NT2RP3000919, highly similar to Rattus norvegicus golgi peripheral membrane protein p65 mRNA. /FEA=mRNA /DB_XREF=gi:7022910 /UG=Hs.4291 hypothetical protein FLJ23443	AK001574		NP_114105
215758_x_at	0.041055	HTF34-like ZNF gene; Homo sapiens chromosome 19, BAC 273239 (CIT-B-320G13), complete sequence.	AC007204		
215780_s_at	0.003539	Human DNA sequence from PAC 30P20 on chromosome Xq21.1-Xq21.3. Contains set pseudogene, ESTs and STS.	Z95126		

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215806_x_at	0.026891	Consensus includes gb:M13231.1 /DEF=Human T-cell receptor aberrantly rearranged gamma-chain mRNA from cell line HPB-MLT. /FEA=mRNA /DB_XREF=gi:339168 /UG=Hs.274509 T cell receptor gamma constant 2	M13231		
215820_x_at	0.031948	Consensus includes gb:AK001861.1 /DEF=Homo sapiens cDNA FLJ10999 fis, clone PLACE1002529, highly similar to Homo sapiens mRNA for KIAA0713 protein. /FEA=mRNA /DB_XREF=gi:7023392 /UG=Hs.283881 KIAA0713 protein	AK001861		NP_055947
215832_x_at	3.26E-04	ubiquitin specific protease 2	AV722190	Hs.199160	NP_009097
215854_at	0.042466	ESTs	AU146050	Hs.287459	
215883_at	0.010205	Consensus includes gb:AK022326.1 /DEF=Homo sapiens cDNA FLJ12264 fis, clone MAMMA1001600. /FEA=mRNA /DB_XREF=gi:10433699 /UG=Hs.288481 Homo sapiens cDNA FLJ12264 fis, clone MAMMA1001600	AK022326		
215884_s_at	0.016934	Consensus includes gb:AK001029.1 /DEF=Homo sapiens cDNA FLJ10167 fis, clone HEMBA1003617, highly similar to Homo sapiens HRIHFB2157 mRNA. /FEA=mRNA /DB_XREF=gi:7022050 /UG=Hs.4552 ubiquitin-2	AK001029		NP_038472
215888_at	6.68E-04	Consensus includes gb:AK026889.1 /DEF=Homo sapiens cDNA: FLJ23236 fis, clone COL00725. /FEA=mRNA /DB_XREF=gi:10439854 /UG=Hs.288786 Homo sapiens cDNA: FLJ23236 fis, clone COL00725	AK026889		

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215930_s_at	0.044231	Consensus includes gb:U73682.1 /DEF=Human meningioma-expressed antigen 11 (MEA11) mRNA, partial cds. /FEA=mRNA /GEN=MEA11 /PROD=meningioma-expressed antigen 11 /DB_XREF=gi:1916671 /UG=Hs.117242 meningioma expressed antigen 6 (coiled-coil proline-rich)	U73682		NP_005921
215933_s_at	2.05E-04	Consensus includes gb:Z21533.1 /DEF=H.sapiens: HEX gene encoding homeobox related protein. /FEA=mRNA /GEN=HEX /PROD=homeobox related protein /DB_XREF=gi:32068 /UG=Hs.118651 hematopoietically expressed homeobox	Z21533		NP_002720
215938_s_at	0.022752	Consensus includes gb:AK001290.1 /DEF=Homo sapiens cDNA FLJ10428 fis, clone NT2RP1000376, highly similar to Homo sapiens mRNA; cDNA DKFZp434A102. /FEA=mRNA /DB_XREF=gi:7022453 /UG=Hs.120360 phospholipase A2, group VI (cytosolic, calcium-independent)	AK001290		
215952_s_at	0.041055	Consensus includes gb:AF090094.1 /DEF=Homo sapiens clone IMAGE 172979. /FEA=mRNA /DB_XREF=gi:4063629 /UG=Hs.125078 ornithine decarboxylase antizyme 1	AF090094		
215963_x_at	1.26E-04	Human DNA sequence from clone RP1-111B22 on chromosome 6q16-21, complete sequence.	Z98200		
215966_x_at	8.68E-04	glycerol kinase	AA292874	Hs.1466	NP_000158
215967_s_at	0.019657	lymphocyte antigen 9	AL582804	Hs.153042	NP_002339

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215977_x_at	0.016934	Consensus includes gb:X68285.1 /DEF=H.sapiens mRNA for glycerol kinase. /FEA=mRNA /PROD=glycerol kinase /DB_XREF=gi:38413 /UG=Hs.1466 glycerol kinase	X68285		NP_000158
215978_x_at	0.001116	Consensus includes gb:AK021514.1 /DEF=Homo sapiens cDNA FLJ11452 fis, clone HEMBA1001435. /FEA=mRNA /DB_XREF=gi:10432710 /UG=Hs.148598 Homo sapiens cDNA FLJ11452 fis, clone HEMBA1001435	AK021514		
215983_s_at	0.023856	Consensus includes gb:D83768.1 /DEF=Human clone N9S Rep-8 mRNA, partial cds. /FEA=mRNA /GEN=Rep-8 /DB_XREF=gi:1913786 /UG=Hs.153678 reproduction 8	D83768		NP_005662
215990_s_at	0.003974	Consensus includes gb:S67779.1 /DEF=BCL5=Cys2-His2 zinc-finger transcription factor human, liver, mRNA, 2600 nt. /FEA=mRNA /DB_XREF=gi:459372 /UG=Hs.155024 B-cell CLL lymphoma 6 (zinc finger protein 51)	S67779		
216012_at	0.041055	Consensus includes gb:U43604.1 /DEF=Human unidentified mRNA, partial sequence. /FEA=mRNA /DB_XREF=gi:1171236 /UG=Hs.159901 Human unidentified mRNA, partial sequence	U43604		
216033_s_at	0.003171	Consensus includes gb:S74774.1 /DEF=p59fyn(T)=OKT3-induced calcium influx regulator human, Jurkat J6 T cell line, mRNA Partial, 1605 nt. /FEA=CDS /PROD=tyrosine kinase p59fyn(T) /DB_XREF=gi:802050 /UG=Hs.169370 FYN oncogene related to SRC, FGR, YES	S74774		NP_694593

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216035_x_at	5.64E-04	transcription factor 7-like 2 (T-cell specific, HMG-box)	AV721430	Hs.348412	NP_110383
216037_x_at	0.00228	transcription factor 7-like 2 (T-cell specific, HMG-box)	AA664011	Hs.348412	NP_110383
216041_x_at	0.041055	Consensus includes gb:AK023348.1 /DEF=Homo sapiens cDNA FLJ13286 fis, clone OVARC1001154, highly similar to Homo sapiens clone 24720 epithelin 1 and 2 mRNA. /FEA=mRNA /DB_XREF=gi:10435243 /UG=Hs.180577 granulin	AK023348		NP_002078
216054_x_at	5.03E-04	Human MLC1emb gene for embryonic myosin alkaline light chain, promoter and exon 1.	NM_002476		NP_002467
216063_at	0.001477	ESTs, Weakly similar to HBE_HUMAN HEMOGLOBIN EPSILON CHAIN [H.sapiens]	N55205	Hs.20205	
216071_x_at	0.010205	intelligence reducing insertion protein INGRIN; Homo sapiens OPA-containing protein (HOPA) gene, complete cds.	NM_005120		NP_005111
216088_s_at	0.00228	Consensus includes gb:AL078633 /DEF=Human DNA sequence from clone RP5-1005F21 on chromosome 20 Contains three isoforms of the gene PSMA7 (proteasome (prosome, macropain) subunit, alpha type, 7), the gene HRH3 encoding the Histamine H3 receptor, the gene KIAA0693 (similar to SSXT, sy... /FEA=CDS_4 /DB_XREF=gi:9408092 /UG=Hs.233952 proteasome (prosome, macropain) subunit, alpha type, 7	AL078633		
216090_x_at	0.023856	Consensus includes gb:AL078636.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 117929. /FEA=mRNA /DB_XREF=gi:5051866 /UG=Hs.227171 Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 117929	AL078636		

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
216091_s_at	0.016934	Consensus includes gb:AF101784.1 /DEF=Homo sapiens b-TRCP variant E3RS-lkappaB mRNA, partial cds. /FEA=mRNA /PROD=b-TRCP variant E3RS-lkappaB /DB_XREF=gi:4165135 /UG=Hs.226434 beta-transducin repeat containing	AF101784		NP_378663
216095_x_at	0.00228	Consensus includes gb:AF057354.1 /DEF=Homo sapiens myotubularin-related protein 1a mRNA, partial cds. /FEA=mRNA /PROD=myotubularin-related protein 1a /DB_XREF=gi:5138901 /UG=Hs.23200 myotubularin related protein 1	AF057354		NP_789746
216100_s_at	6.79E-04	DKFZP586G011:protein	BG289527	Hs.234265	NP_056417
216105_x_at	0.03002	Consensus includes gb:X86428 /DEF=Homo sapiens PTPA gene for phosphotyrosyl phosphatase activator, exon 1 and joined CDS /FEA=mRNA_2 /DB_XREF=gi:974836 /UG=Hs.236963 protein phosphatase 2A, regulatory subunit B (PR 53)	X86428		
216110_x_at	0.046846	Consensus includes gb:AU147017 /FEA=EST /DB_XREF=gi:11008538 /DB_XREF=est:AU147017 /CLONE=HEMBB1002152 /UG=Hs.301905 Homo sapiens cDNA FLJ14080 fis, clone HEMBB1002152	AK024142		
216112_at	0.028893	AU157200 PLACE1 Homo sapiens cDNA clone PLACE1006617 3', mRNA sequence.	AU157200		
216125_s_at	0.001323	Consensus includes gb:AF064606.1 /DEF=Homo sapiens KB07 protein mRNA, partial cds. /FEA=mRNA /PROD=KB07 protein /DB_XREF=gi:3152665 /UG=Hs.306242 Homo sapiens KB07 protein mRNA, partial cds	AF064606		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216126_at	0.022752	Consensus includes gb:AK022418.1 /DEF=Homo sapiens cDNA FLJ12356 fis, clone MAMMA1002339. /FEA=mRNA /DB_XREF=gi:10433811 /UG=Hs.306635 Homo sapiens cDNA FLJ12356 fis, clone MAMMA1002339	AK022418		
216187_x_at	0.013223	Consensus includes gb:AF222691.1 /DEF=Homo sapiens Alu repeat (LNX1) mRNA sequence. /FEA=mRNA /DB_XREF=gi:12655850 /UG=Hs.307008 Homo sapiens Alu repeat (LNX1) mRNA sequence	AF222691		
216191_s_at	0.004341	Consensus includes gb:X72501.1 /DEF=H.sapiens mRNA for soluble delta TCR. /FEA=CDS /PROD=gamma-delta T-cell receptor /DB_XREF=gi:298104 /UG=Hs.307187 H.sapiens mRNA for soluble delta TCR	X72501		
216202_s_at	0.016686	Consensus includes gb:U15555.1 /DEF=Human serine palmitoyltransferase (LCB2) mRNA, partial cds. /FEA=mRNA /GEN=LCB2 /PROD=serine palmitoyltransferase /DB_XREF=gi:1001944 /UG=Hs.59403 serine palmitoyltransferase, long chain base subunit 2	U15555		NP_004854
216207_x_at	0.045316	immunoglobulin kappa constant	AW408194	Hs.306357	
216218_s_at	1.26E-04	Consensus includes gb:AK023546.1 /DEF=Homo sapiens cDNA FLJ13484 fis, clone PLACE1003888, weakly similar to 1-PHOSPHATIDYLINOSITOL-4,5-BISPHOSPHATE PHOSPHODIESTERASE DELTA 1 (EC 3.1.4.11). /FEA=mRNA /DB_XREF=gi:10435512 /UG=Hs.54886 phospholipase C, epsilon 2	AK023546		NP_055999

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216221_s_at	7.58E-04	Consensus includes gb:D87078.2 /DEF=Homo sapiens mRNA for KIAA0235 protein, partial cds. /FEA=mRNA /GEN=KIAA0235 /PROD=KIAA0235 protein /DB_XREF=gi:6634004 /UG=Hs.6151 pumilio (Drosophila) homolog 2	D87078		NP_056132
216266_s_at	0.026891	Consensus includes gb:AK025637.1 /DEF=Homo sapiens cDNA: FLJ21984 fis, clone HEP06222, highly similar to AF111162 Homo sapiens guanine nucleotide exchange factor mRNA. /FEA=mRNA /DB_XREF=gi:10438216 /UG=Hs.94631 brefeldin A-inhibited guanine nucleotide-exchange protein 1	AK025637		NP_006412
216268_s_at	1.28E-04	Consensus includes gb:U77914.1 /DEF=Human soluble protein Jagged mRNA, partial cds. /FEA=mRNA /PROD=soluble protein Jagged /DB_XREF=gi:1684889 /UG=Hs.91143 jagged 1 (Alagille syndrome)	U77914		NP_000205
216274_s_at	3.26E-04	signal peptidase complex (18kD)	N99438	Hs.9534	NP_055115
216282_x_at	0.016934	Consensus includes gb:AJ224143.1 /DEF=Homo sapiens mRNA variant alfa for RNA polymerase II subunit 3. /FEA=mRNA /GEN=RPB3 /PROD=RNA polymerase II subunit /DB_XREF=gi:2920708 /UG=Hs.79402 polymerase (RNA) II (DNA directed) polypeptide C (33kD)	AJ224143		NP_116558
216288_at	6.68E-04	AU159276 THYRO1 Homo sapiens cDNA clone THYRO1001262 3', mRNA sequence.	AU159276		NP_006630
216295_s_at	0.010205	Consensus includes gb:X81636.1 /DEF=H.sapiens clathrin light chain a gene. /FEA=mRNA /DB_XREF=gi:704460 /UG=Hs.285688 H.sapiens clathrin light chain a gene	X81636		

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
216296_at	0.022988	Consensus includes gb:X81636.1 /DEF=H.sapiens clathrin light chain a gene. /FEA=mRNA /DB_XREF=gi:704460 /UG=Hs.285688 H.sapiens clathrin light chain a gene	X81636		
216304_x_at	0.041055	Homo sapiens FTSH gene for putative ATPases, exons 1 and 2 and join CDS.	AJ295618		
216308_x_at	0.007779	Consensus includes gb:AK026752.1 /DEF=Homo sapiens cDNA: FLJ23099 fis; clone LNG07460, highly similar to AF113251 Homo sapiens putative 2-hydroxyacid dehydrogenase mRNA. /FEA=mRNA /DB_XREF=gi:10439675 /UG=Hs.155742 glyoxylate reductasehydroxypyruvate reductase	AK026752		NP_036335
216321_s_at	0.00161	Consensus includes gb:X03348.1 /DEF=Human mRNA for beta-glucocorticoid receptor (clone OB10). /FEA=mRNA /PROD=beta-glucocorticoid receptor /DB_XREF=gi:31681 /UG=Hs.75772 nuclear receptor subfamily 3, group C, member 1	X03348		NP_000167
216323_x_at	0.001303	alpha-tubulin isotype H2-alpha; Human alpha-tubulin isotype H2-alpha gene, last exon.	K03460		
216379_x_at	3.26E-04	Consensus includes gb:AK000168.1 /DEF=Homo sapiens cDNA FLJ20161 fis; clone COL09252, highly similar to L33930 Homo sapiens CD24 signal transducer mRNA. /FEA=mRNA /DB_XREF=gi:7020079 /UG=Hs.332045 Homo sapiens cDNA FLJ20161 fis; clone COL09252, highly similar to L33930 Homo sapiens CD24 signal transducer mRNA	AK000168		

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
216383_at	2.05E-04	Homo sapiens chromosome X clone Qc-7G6, QLL-F1720, QLL-C1335, Qc-8B7, Qc-11H12, Qc-7F6, QLL-E153, Qc-10E8, Qc-10B7 map q28, complete sequence.	U52111		
216388_s_at	0.00884	R2, Human putative G-protein-coupled receptor (GPR16) gene, complete cds.	U33448		
216389_s_at	0.021312	Consensus includes gb:AF283773.2 /DEF=Homo Sapiens clone TCBAP0888 mRNA sequence. /FEA=mRNA /DB_XREF=gi:10281743 /UG=Hs.283976 hypthetical protein PRO2389	AF283773		NP_079506
216409_at	0.001116	Consensus includes gb:AL390168.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547D104 (from clone DKFZp547D104). /FEA=mRNA /GEN=DKFZp547D104 /PROD=hypothetical protein /DB_XREF=gi:9368863 /UG=Hs.287371 hypothetical protein DKFZp547D104	AL390168		
216443_at	0.034317	Consensus includes gb:AK024615.1 /DEF=Homo sapiens cDNA: FLJ20962 fis, clone ADSh00804. /FEA=mRNA /DB_XREF=gi:10436931 /UG=Hs.306698 Homo sapiens cDNA: FLJ20962 fis, clone ADSh00804	AK024615		
216457_s_at	3.26E-04	Consensus includes gb:AK026080.1 /DEF=Homo sapiens cDNA: FLJ22427 fis, clone HRC09013. /FEA=mRNA /DB_XREF=gi:10438814 /UG=Hs.288883 splicing factor 3a, subunit 1, 120kD	AK026080		
216483_s_at	0.020793	Hypothetical partial human protein; Homo sapiens chromosome 19, cosmid R33729, complete sequence.	AC005339		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216484_x_at	1.26E-04	Consensus includes gb:L24521.1 /DEF=Human transformation-related protein mRNA, 3 end. /FEA=mRNA /PROD=transformation-related protein /DB_XREF=gi:403459 /UG=Hs.300705 Human transformation-related protein mRNA, 3 end	L24521		NP_004485
216508_x_at	0.016934	similar to nonhistone chromosomal protein HMG-1 [Homo sapiens]; probable pseudogene; similar to P09429 (PID:g123369); H_NH0244E06.1; Homo sapiens BAC clone RP11-244E6 from 2, complete sequence.	AC007277		
216511_s_at	0.005852	Consensus includes gb:AJ270770 /DEF=Homo sapiens partial TCF-4 gene for T-cell transcription factor-4, exon 1 and joined CDS features /FEA=mRNA_1 /DB_XREF=gi:9188625 /UG=Hs.283857 Homo sapiens partial TCF-4 gene for T-cell transcription factor-4, exon 1 and joined CDS features	AJ270770		
216520_s_at	0.005852	Homo sapiens HDCMB21P gene, complete cds.	AF072098		
216526_x_at	0.041055	Consensus includes gb:AK024836.1 /DEF=Homo sapiens cDNA: FLJ21183 fis, clone CAS11634, highly similar to HSHLACW07 Homo sapiens mRNA for human leukocyte antigen C alpha chain. /FEA=mRNA /DB_XREF=gi:10437242 /UG=Hs.277477 major histocompatibility complex, class I, C	AK024836		NP_002108

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
216550_x_at	0.026891	Consensus includes gb:X80821.1 /DEF=H.sapiens mRNA for ribosomal protein L18a homologue. /FEA=mRNA /PROD=ribosomal protein L18a homologue /DB_XREF=gi:527579 /UG=Hs.302177 H.sapiens mRNA for ribosomal protein L18a homologue	X80821		NP_056023
216556_x_at	0.021197	Consensus includes gb:AL135926 /DEF=Human DNA sequence from clone RP11-375F2 on chromosome 1 Contains a pseudogene similar to UBL1 (ubiquitin-like 1 (sentrin)), a pseudogene similar to ribosomal protein L29, ESTs, STSs and GSSs /FEA=CDS_1 /DB_XREF=gi:9801286. /UG=Hs.302113 Human DNA sequence from clone RP11-375F2 on chromosome 1 Contains a pseudogene similar to UBL1 (ubiquitin-like 1 (sentrin)), a pseudogene similar to ribosomal protein L29, ESTs, STSs and GSSs	AL135926		
216563_at	6.68E-04	Consensus includes gb:X80821.1 /DEF=H.sapiens mRNA for ribosomal protein L18a homologue. /FEA=mRNA /PROD=ribosomal protein L18a homologue /DB_XREF=gi:527579 /UG=Hs.302177 H.sapiens mRNA for ribosomal protein L18a homologue	X80821		NP_056023

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
216565_x_at	0.016934	dJ781L3.1 (similar to IFITM3 (interferon induced transmembrane protein 3 (1-8U))) match: proteins: Sw:Q01628 Sw:Q01629 Tr:Q9R175 Tr:Q9R176 Sw:Q91499 Sw:P26376 Tr:O88728 Sw:P13164; Human DNA sequence from clone RP4-781L3 on chromosome 1p34.3-36.11 Contains a pseudogene similar to IFITM3 (interferon induced transmembrane protein 3 (1-8U)), STSs and GSSs, complete sequence.	AL121994		
216570_x_at	0.004341	Consensus includes gb:AL096829 /DEF=Human DNA sequence from clone RP4-595K12 on chromosome 1p31.2-31.3 Contains a pseudogene similar to 60S RPL29 (ribosomal protein L29 (cell surface heparin binding protein HIP)), a chromosome 1 specific mRNA (KIAA0499), a novel mRNA (KIAA0433), EST... /FEA=CDS /DB_XREF=gi:6634461 /UG=Hs.302120 Human DNA sequence from clone RP4-595K12 on chromosome 1p31.2-31.3 Contains a pseudogene similar to 60S RPL29 (ribosomal protein L29 (cell surface heparin binding protein HIP)), a chromosome 1 specific mRNA (KIAA0499), a novel mRNA (KIAA0433), ESTs, STSs,	AL096829		
216593_s_at	0.003171	pseudogene of PIGC; see also AB000360 putative; Homo sapiens PIGCP1 pseudogene.	AB000359		

Gene Identifier	p-value	Description	Gene Accession No.	Unig n Acc ssion No.	Protein Acc ssion No.
216606_x_at	0.029453	Consensus includes gb:AL050332 /DEF=Human DNA sequence from clone RP4-570F3 on chromosome 6. Contains a gene similar to Rattus norvegicus synaptic ras GTPase-activating protein p135, the CICK0721Q.5 (polypeptide from patented cDNA Em:E06811) gene, the PHF1 (PHD finger protein 1) gen... /FEA=mRNA_1 /DB_XREF=gi:6010176 /UG=Hs.306238 acyl-protein	AL050332		
216640_s_at	0.041055	Consensus includes gb:AK026926.1 /DEF=Homo sapiens cDNA: FLJ23273 fis, clone HEP02611, highly similar to HSU79278 Human protein disulfide isomerase-related protein P5 mRNA. /FEA=mRNA /DB_XREF=gi:10439898 /UG=Hs.182429 protein disulfide isomerase-related protein	AK026926		
216652_s_at	1.26E-04	Consensus includes gb:AL137673.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434H0872 (from clone DKFZp434H0872). /FEA=mRNA /DB_XREF=gi:6807841 /UG=Hs.306454 Homo sapiens mRNA; cDNA DKFZp434H0872 (from clone DKFZp434H0872)	AL137673		
216739_at	0.009716	Consensus includes gb:AK024527.1 /DEF=Homo sapiens cDNA: FLJ20874 fis, clone ADKA02818. /FEA=mRNA /DB_XREF=gi:10436829 /UG=Hs.306684 Homo sapiens cDNA: FLJ20874 fis, clone ADKA02818	AK024527		
216751_at	0.02145	Consensus includes gb:AK024879.1 /DEF=Homo sapiens cDNA: FLJ21226 fis, clone COL00721. /FEA=mRNA /DB_XREF=gi:10437291 /UG=Hs.306715 Homo sapiens cDNA: FLJ21226 fis, clone COL00721	AK024879		

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
216756_at	8.68E-04	Consensus includes gb:AK024995.1 /DEF=Homo sapiens cDNA: FLJ21342 fis, clone COL02673. /FEA=mRNA /DB_XREF=gi:10437426 /UG=Hs.306739 Homo sapiens cDNA: FLJ21342 fis, clone COL02673	AK024995		
216791_at	0.006096	Consensus includes gb:AK026449.1 /DEF=Homo sapiens cDNA: FLJ22796 fis, clone KAIA2544. /FEA=mRNA /DB_XREF=gi:10439318 /UG=Hs.306850 Homo sapiens cDNA: FLJ22796 fis, clone KAIA2544	AK026449		
216804_s_at	0.011725	Consensus includes gb:AK027217.1 /DEF=Homo sapiens cDNA: FLJ23564 fis, clone LNG10773. /FEA=mRNA /DB_XREF=gi:10440296 /UG=Hs.306913 Homo sapiens cDNA: FLJ23564 fis, clone LNG10773	AK027217		
216833_x_at	0.034317	Consensus includes gb:U05255.1 /DEF=Human glycoporphin HeP2 mRNA, partial cds. /FEA=mRNA /PROD=glycoporphin HeP2 /DB_XREF=gi:454085 /UG=Hs.307185 Human glycoporphin HeP2 mRNA, partial cds	U05255		NP_002091
216835_s_at	2.05E-04	Consensus includes gb:AF035299.1 /DEF=Homo sapiens clone 23863 mRNA, partial cds. /FEA=mRNA /DB_XREF=gi:2661055 /UG=Hs.103854 docking protein 1, 62kD (downstream of tyrosine kinase 1)	AF035299		NP_001372
216841_s_at	7.58E-04	Consensus includes gb:X15132.1 /DEF=Human mRNA for manganese containing superoxide dismutase (EC 1.15.1.1). /FEA=mRNA /DB_XREF=gi:34794 /UG=Hs.318885 superoxide dismutase 2, mitochondrial	X15132		NP_000627

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216858_x_at	0.018973	Consensus includes gb:AL080112.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H0722 (from clone DKFZp586H0722). /FEA=mRNA /DB_XREF=gi:5262539 /UG=Hs.332731 Homo sapiens mRNA; cDNA DKFZp586H0722 (from clone DKFZp586H0722)	AL080112		
216870_x_at	0.007779	Consensus includes gb:AF264787.1 /DEF=Homo sapiens BCMS-upstream neighbor (BCMSUN) mRNA, partial sequence. /FEA=mRNA /DB_XREF=gi:11141510 /UG=Hs.43628 deleted in lymphocytic leukemia, 2	AF264787		NP_006012
216899_s_at	0.007779	Homo sapiens PAC clone RP5-1139P1 from 7p11.2-p21, complete sequence.	AC003999		
216902_s_at	0.009716	Consensus includes gb:AF001549 /DEF=Human Chromosome 16 BAC clone CIT987SK-A-270G1 /FEA=mRNA_1 /DB_XREF=gi:3355302 /UG=Hs.110103 RNA polymerase I transcription factor RRN3	AF001549		
216903_s_at	1.32E-04	Consensus includes gb:AK022697.1 /DEF=Homo sapiens cDNA FLJ12635 fis, clone NT2RM4001865, highly similar to Homo sapiens mRNA for atopy related autoantigen CALC. /FEA=mRNA /DB_XREF=gi:10434244 /UG=Hs.61628 calcium binding atopy-related autoantigen 1	AK022697		NP_006068
216942_s_at	1.26E-04	Consensus includes gb:D28586.1 /DEF=Human mRNA for LFA-3(delta D2); partial cds. /FEA=mRNA /PROD=LFA-3(delta D2) /DB_XREF=gi:466540 /UG=Hs.75626 CD58 antigen, (lymphocyte function-associated antigen 3)	D28586		NP_001770

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
216950_s_at	0.003171	Consensus includes gb:X14355.1 /DEF=Human mRNA for high affinity Fc receptor (FcRI) b form. /FEA=mRNA /DB_XREF=gi:31333 /UG=Hs.77424 Fc fragment of IgG, high affinity Ia, receptor for (CD64)	X14355		
216988_s_at	2.05E-04	Homo sapiens (clone hh18) protein tyrosine phosphatase (ptp-IV1r) gene, 5' end of cds.	L48722		
216993_s_at	0.016686	Consensus includes gb:U32169 /DEF=Human pro-a2 chain of collagen type XI (COL11A2) gene, complete cds /FEA=mRNA_3 /DB_XREF=gi:1000744 /UG=Hs.121509 collagen, type XI, alpha 2	U32169		
217003_s_at	0.041795	Consensus includes gb:AJ132823.1 /DEF=Homo sapiens mRNA for tMDC II, isoform d. /FEA=mRNA /DB_XREF=gi:4468348 /UG=Hs.194768 a disintegrin and metalloproteinase domain 5	AJ132823		
217009_at	0.018443	Consensus includes gb:AL121974 /DEF=Human DNA sequence from clone RP3-417L20 on chromosome 6p12-21.3. Contains the 5 end of the gene for cysteine-rich secretory protein 3 (CRISP-3, SGP28), the PGK2 gene for phosphoglycerate kinase 2, STSs and GSSs /FEA=CDS /DB_XREF=gi:9828129 /UG=Hs.78771 phosphoglycerate kinase 1	AL121974		
217016_x_at	0.003171	Consensus includes gb:AK026825.1 /DEF=Homo sapiens cDNA: FLJ23172 fis, clone LNG10005. /FEA=mRNA /DB_XREF=gi:10439771 /UG=Hs.306885 Homo sapiens cDNA: FLJ23172 fis, clone LNG10005	AK026825		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217041_at	0.017634	Consensus includes gb:AL162057.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761N0524 (from clone DKFZp761N0524). /FEA=mRNA /DB_XREF=gi:7328105 /UG=Hs.91622 neuronal pentraxin receptor	AL162057		NP_478058
217042_at	0.003171	Consensus includes gb:AL096716.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564M1462 (from clone DKFZp564M1462); partial cds. /FEA=mRNA /GEN=DKFZp564M1462 /PROD=hypothetical protein /DB_XREF=gi:5419850 /UG=Hs.226007 DKFZP564M1462 protein	AL096716		
217043_s_at	1.26E-04	Consensus includes gb:U95822.1 /DEF=Human putative transmembrane GTPase mRNA, partial cds. /FEA=mRNA /GEN=fzo /PROD=putative transmembrane GTPase /DB_XREF=gi:2252803 /UG=Hs.197877 hypothetical protein FLJ20693	U95822		
217078_s_at	0.003597	Consensus includes gb:AJ010102.1 /DEF=Homo Sapiens mRNA for Natural killer cell IRC1c gene. /FEA=mRNA /GEN=IRC1c /DB_XREF=gi:5706457 /UG=Hs.9688 leukocyte membrane antigen	AJ010102		
217118_s_at	2.05E-04	Consensus includes gb:AK025608.1 /DEF=Homo sapiens cDNA: FLJ21955 fis, clone HEP05265. /FEA=mRNA /DB_XREF=gi:10438177 /UG=Hs.13255 KIAA0930 protein	AK025608		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217122_s_at	0.033381	Consensus includes gb:AL031282 /DEF=Human DNA sequence from clone 283E3 on chromosome 1p36.21-36.33. Contains the alternatively spliced gene for Matrix Metalloproteinase in the Female Reproductive tract MIFR1, -2, MMP2122A, -B and -C, a novel gene, the alternatively spliced CDC2L2 ... /FEA=mRNA_6 /DB_XREF=gi:3860395 /UG=Hs.214646 KIAA0447 gene product	AL031282		
217125_at	0.005692	Homo sapiens UBBP2 pseudogene for ubiquitin UBB.	X04802		
217143_s_at	0.02145	Consensus includes gb:X06557.1 /DEF=Human mRNA for TCR-delta chain. /FEA=mRNA /DB_XREF=gi:37003 /UG=Hs.2014 T cell receptor delta locus	X06557		
217144_at	1.28E-04	Homo sapiens UBBP1 pseudogene for ubiquitin UBB.	X04801		
217164_at	0.010205	Consensus includes gb:AK024108.1 /DEF=Homo sapiens cDNA FLJ14046 fis, clone HEMBA1006461. /FEA=mRNA /DB_XREF=gi:10436406 /UG=Hs.142677 Homo sapiens cDNA FLJ14046 fis, clone HEMBA1006461	AK024108		
217167_x_at	0.003862	Consensus includes gb:AJ252550 /DEF=Homo sapiens partial GK gene for glycerol kinase, exon 1 (glycerol kinase deficiency case) /FEA=CDS_1 /DB_XREF=gi:5834425 /UG=Hs.1466 glycerol kinase	AJ252550		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217168_s_at	0.026891	Consensus includes gb:AF217990.1 /DEF=Homo sapiens clone PP1722 unknown mRNA. /FEA=mRNA /PROD=unknown /DB_XREF=gi:10441910 /UG=Hs.146393 homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1	AF217990		NP_055500
217200_x_at	7.58E-04	Consensus includes gb:U06715.1 /DEF=Human cytochrome B561, HCYTO B561, mRNA, partial cds. /FEA=mRNA /GEN=B561 /PROD=HCYTO B561 /DB_XREF=gi:476590 /UG=Hs.153028 cytochrome b-561	U06715		
217208_s_at	0.020576	match: proteins: Tr:Q62402 Sw:Q62696 Sw:Q12959; Human DNA sequence from clone RP5-1061C18 on chromosome 3p36.22-36.33 Contains the 5' part of the DLG1 gene encoding the discs protein (large Drosophila homolog 1, presynaptic protein SAP97) and a CpG island, complete sequence.	AL121981		
217209_at	0.005673	Human gene for carcinoembryonic antigen subdomains A and B.	X16454		
217211_at	6.79E-04	Human beta-cytoplasmic actin (ACTBP9) pseudogene.	D50604		
217212_s_at	7.44E-04	Human DNA sequence from phage LAW2 from a contig from the tip of the short arm of chromosome 16, spanning 2Mb of 16p13.3 Contains Interleukin 9 receptor pseudogene.	Z84723		
217226_s_at	0.026891	Consensus includes gb:M95929.1 /DEF=Human homeobox protein (PHOX1) mRNA, 3 end. /FEA=mRNA /GEN=PHOX1 /PROD=homeobox protein /DB_XREF=gi:189946 /UG=Hs.155606 paired mesoderm homeo box 1	M95929		
217232_x_at	0.001116	Homo sapiens mutant beta-globin (HBB) gene, complete cds.	AF059180		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217249_x_at	0.005852	cytochrome C oxidase; match to P14406 (PID:g117121); H_RG162B04.1; Homo sapiens BAC clone CTB-162B4 from 4, complete sequence.	AC004544		
217274_x_at	0.001205	Consensus includes gb:X52005.1 /DEF=H.sapiens skeletal embryonic myosin light chain 1 (MLC1) mRNA. /FEA=mRNA /GEN=MCL1 /PROD=myosin light chain 1 /DB_XREF=gi:34677 /UG=Hs.159218 H.sapiens skeletal embryonic myosin light chain 1 (MLC1) mRNA	X52005		
217286_s_at	0.007779	Consensus includes gb:BC001805.1 /DEF=Homo sapiens, clone IMAGE:3543670, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3543670) /DB_XREF=gi:12804742 /UG=Hs.240615 hypothetical protein FLJ13556 similar to N-myc downstream regulated 3	BC001805		
217294_s_at	0.029453	Consensus includes gb:U88968.1 /DEF=Human alpha enolase like 1 (ENO1L1) mRNA, partial cds. /FEA=mRNA /GEN=ENO1L1 /PROD=alpha enolase like 1 /DB_XREF=gi:3282242 /UG=Hs.254105 enolase 1, (alpha)	U88968		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217356_s_at	0.033381	Consensus includes gb:S81916.1 /DEF=phosphoglycerate kinase {alternatively spliced} human, phosphoglycerate kinase deficient patient with episodes of muscle, mRNA Partial Mutant, 307 nt. /FEA=mRNA /PROD=phosphoglycerate kinase /DB_XREF=gi:1470308 /UG=Hs.169313 Phosphoglycerate kinase {alternatively spliced} human, phosphoglycerate kinase deficient patient with episodes of muscle, mRNA Partial Mutant, 307 nt	S81916		
217362_x_at	0.001551	Consensus includes gb:AF005487.1 /DEF=Homo sapiens MHC class II antigen (DRB6) mRNA, HLA-DRB6*0201 allele, sequence. /FEA=mRNA /DB_XREF=gi:5915893 /UG=Hs.167385 Homo sapiens MHC class II antigen HLA-DRB6 mRNA, partial cds	AF005487		
217371_s_at	0.010644	Consensus includes gb:Y09908.1 /DEF=H.sapiens mRNA for interleukin-15. /FEA=mRNA /GEN=IL-15 /PROD=interleukin-15 /DB_XREF=gi:2143255 /UG=Hs.168132 interleukin 15	Y09908		
217393_x_at	0.021165	match: proteins: Sw:P52490 Sw:P35128 Sw:Q29503 Sw:P52492 Tr:Q42045 Tr:Q9ZVA6 Tr:Q9ZTZ8 Tr:Q42897 Sw:P35131 Sw:P35132 Sw:P35133 Sw:P35134 Sw:P35135 Wp:CE19850 Tr:O13685 Tr:Q43780 Sw:Q16781 Tr:O94098; Human DNA sequence from clone RP3-526F5 on chromosome Xq26.3-28. Contains the gene for a novel protein similar to UBE2N (ubiquitin-conjugating enzyme E2N (homologous to yeast UBC13)), ESTs, STSs and GSSs, complete sequence.	AL109622		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217394_at	0.026891	Consensus includes gb:AE000659 /DEF=Homo sapiens T-cell receptor alpha delta locus from bases 250472 to 501670 (section 2 of 5) of the Complete Nucleotide Sequence /FEA=CDS_9 /DB_XREF=gi:2358025 /UG=Hs.272547 Homo sapiens T-cell receptor alpha delta locus from bases 250472 to 501670 (section 2 of 5) of the Complete Nucleotide Sequence	AE000659		
217403_s_at	0.041795	Consensus includes gb:AC074331 /DEF=Homo sapiens chromosome 19, BAC CIT-HSPC_204F22 (BC228680), complete sequence; contains bacterial insertion element /FEA=CDS_4 /DB_XREF=gi:9502399 /UG=Hs.9615 myosin regulatory light chain 2, smooth muscle isoform	AC074331		
217414_x_at	3.56E-04	Human alpha-globin gene with flanks.	NM_000517		NP_000508
217418_x_at	0.018693	Consensus includes gb:X12530.1 /DEF=Human mRNA for B lymphocyte antigen CD20 (B1, Bp35). /FEA=mRNA /DB_XREF=gi:29773 /UG=Hs.89751 membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE, high affinity I, receptor for; beta polypeptide)	X12530		NP_690606
217427_s_at	4.42E-04	Consensus includes gb:X75296.1 /DEF=H.sapiens TUP1-like enhancer of split gene 1 (TUPLE1) mRNA. /FEA=mRNA /GEN=TUPLE1 /PROD=TUP1 like enhancer of SPLIT gene 1 /DB_XREF=gi:434982 /UG=Hs.172350 HIR (histone cell cycle regulation defective, S. cerevisiae) homolog A	X75296		NP_003316

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
217437_s_at	2.05E-04	Consensus includes gb:AB029026.1 /DEF=Homo sapiens mRNA for KIAA1103 protein, partial cds. /FEA=mRNA /GEN=KIAA1103 /PROD=KIAA1103 protein /DB_XREF=gi:5689542 /UG=Hs.173159 transforming; acidic coiled-coil containing protein 1	AB029026		NP_006274
217456_x_at	0.00161	Consensus includes gb:M31183.1 /DEF=Homo sapiens (cosmid clone cd3.3) MHC class I HLA-B51 mRNA, 3' end of cds. /FEA=mRNA /GEN=HLA-B /PROD=MHC class I lymphocyte antigen /DB_XREF=gi:601851 /UG=Hs.181392 major histocompatibility complex, class I, E	M31183		NP_005507
217457_s_at	0.028552	Consensus includes gb:X63465.1 /DEF=H.sapiens hGDS mRNA for smg GDS. /FEA=CDS /GEN=hGDS /PROD=smg GDS /DB_XREF=gi:32079 /UG=Hs.7940 RAP1, GTP-GDP dissociation stimulator 1	X63465		NP_066982
217475_s_at	0.028999	Consensus includes gb:AC002073 /DEF=Human PAC clone RP3-515N1 from 22q11.2-q22 /FEA=CDS_1 /DB_XREF=gi:2078469 /UG=Hs.278027 LIM domain kinase 2	AC002073		
217479_at	0.034317	Consensus includes gb:AL110201.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586F1622 (from clone DKFZp586F1622). /FEA=mRNA /DB_XREF=gi:5817120 /UG=Hs.278435 Homo sapiens mRNA; cDNA DKFZp586F1622 (from clone DKFZp586F1622)	AL110201		
217480_x_at	0.005411	Ig kappa chain; Human kappa-immunoglobulin germline pseudogene (cos118) variable region (subgroup V kappa I).	M20812		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217497_at	0.016934	ESTs, Weakly similar to hypothetical protein FLJ20378 [Homo sapiens] [H.sapiens]	AW613387	Hs.388345	NP_001944
217499_x_at	2.17E-04	olfactory receptor, family 7, subfamily E, member 38 pseudogene	AW874308	Hs.409186	
217502_at	0.004341	interferon-induced protein with tetratricopeptide repeats 2	BE888744	Hs.169274	NP_001538
217512_at	0.033491	ESTs, Highly similar to 1312232A kininogen L, high MW [Homo sapiens] [H.sapiens]	BG398937	Hs.289056	NP_000884
217529_at	0.04734	ESTs, Moderately similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	BE547674	Hs.129568	
217534_at	0.015516	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AV720514	Hs.282721	
217540_at	1.26E-04	ESTs, Moderately similar to hypothetical protein FLJ20378 [Homo sapiens] [H.sapiens]	AA721025	Hs.293253	
217552_x_at	3.26E-04	th43e02.x1 NCI_CGAP_Lym12 Homo sapiens cDNA clone IMAGE:2121050 3' similar to contains Alu repetitive element, mRNA:sequence	AI432713		
217560_at	0.02008	ESTs, Weakly similar to hypothetical protein FLJ11267 [Homo sapiens] [H.sapiens]	AI879383	Hs.370291	
217580_x_at	0.041795	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AW301806	Hs.150551	
217610_at	0.042057	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AL047879	Hs.293865	
217614_at	0.02145	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AI439416	Hs.165728	
217627_at	7.58E-04	hypothetical protein FLJ30921	BE515346	Hs.278871	NP_689573
217629_at	0.00228	ESTs, Moderately similar to hypothetical protein FLJ20378 [Homo sapiens] [H.sapiens]	AA365670	Hs.261526	
217673_x_at	0.013223	ESTs, Highly similar to GBAS_HUMAN Guanine nucleotide binding protein G(S), alpha subunit (Adenylate cyclase-stimulating G alpha protein) [H.sapiens]	AA650558	Hs.394484	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217679_x_at	0.005852	tx67h02.x1 NCI_CGAP_Ut1 Homo sapiens cDNA clone IMAGE:2274675 3' similar to contains Alu repetitive element; mRNA sequence.	AI683552		
217682_at	0.004341	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AW503390	Hs.392578	
217695_x_at	0.007934	ESTs, Weakly similar to I57588 HSrel-1 - human (fragment) [H.sapiens]	BF942161	Hs.343402	
217707_x_at	3.26E-04	ESTs	AI535683	Hs.405576	
217713_x_at	0.005039	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AA126763	Hs.182296	
217715_x_at	0.010205	ESTs, Weakly similar to hypothetical protein FLJ20294 [Homo sapiens] [H.sapiens]	BE045142	Hs.274907	
217720_at	0.02145	gb:NM_016139.1 /DEF=Homo sapiens 16.7Kd protein (LOC51142), mRNA. /FEA=mRNA /GEN=LOC51142 /PROD=16.7Kd protein /DB_XREF=gi:7705850 /UG=Hs.180859 16.7Kd protein /FL=gb:BC003079.1 gb:AF078845.1 gb:NM_016139.1	NM_016139		NP_057223
217724_at	0.041055	Consensus includes gb:AF131807.1 /DEF=Homo sapiens clone 25076 mRNA sequence. /FEA=mRNA /DB_XREF=gi:4406639 /UG=Hs.165998 PAI-1 mRNA-binding protein /FL=gb:AL080119.1 gb:NM_015640.1	AF131807		NP_056455
217725_x_at	0.02145	gb:NM_015640.1 /DEF=Homo sapiens PAI-1 mRNA-binding protein (PAI-RBP1), mRNA. /FEA=mRNA /GEN=PAI-RBP1 /PROD=PAI-1 mRNA-binding protein /DB_XREF=gi:7661625 /UG=Hs.165998 PAI-1 mRNA-binding protein /FL=gb:AL080119.1 gb:NM_015640.1	NM_015640		NP_056455

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217727_x_at	1.26E-04	gb:NM_018206.1 /DEF=Homo sapiens vacuolar protein sorting 35 (yeast homolog) (VPS35), mRNA. /FEA=mRNA /GEN=VPS35 /PROD=vacuolar protein sorting 35 (yeast homolog) /DB_XREF=gi:8922645 /UG=Hs.264190 vacuolar protein sorting 35 (yeast homolog) /FL=gb:AF186382.1 gb:AL136888.1 gb:BC002414.1 gb:AF191298.2 gb:NM_018206.1 gb:AF175265.1 gb:AF183418.1	NM_018206		NP_060676
217728_at	3.26E-04	gb:NM_014624.2 /DEF=Homo sapiens S100 calcium-binding protein A6 (calcyclin) (S100A6), mRNA. /FEA=mRNA /GEN=S100A6 /PROD=S100 calcium-binding protein A6 /DB_XREF=gi:9845517 /UG=Hs.275243 S100 calcium-binding protein A6 (calcyclin) /FL=gb:BC001431.1 gb:NM_014624.2	NM_014624		NP_055439
217729_s_at	6.68E-04	gb:NM_001130.3 /DEF=Homo sapiens amino-terminal enhancer of split (AES), mRNA. /FEA=mRNA /GEN=AES /PROD=amino-terminal enhancer of split /DB_XREF=gi:6382065 /UG=Hs.244 amino-terminal enhancer of split /FL=gb:AF072902.1 gb:U04241.1 gb:NM_001130.3	NM_001130		NP_001121
217730_at	0.005852	gb:NM_022152.1 /DEF=Homo sapiens PP1201 protein (PP1201), mRNA. /FEA=mRNA /GEN=PP1201 /PROD=PP1201 protein /DB_XREF=gi:11545897 /UG=Hs.184052 PP1201 protein /FL=gb:NM_022152.1	NM_022152		NP_071435

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
217731_s_at	0.041055	gb:NM_021999.1 /DEF=Homo sapiens integral membrane protein 2B (ITM2B), mRNA. /FEA=mRNA /GEN=ITM2B /PROD=integral membrane protein 2B /DB_XREF=gi:11527401 /UG=Hs.239625 integral membrane protein 2B /FL=gb:NM_021999.1 gb:AF136973.1 gb:BC000554.1 gb:AF092128.1 gb:AF152462.1 gb:AF246221.1	NM_021999		NP_068839
217736_s_at	5.03E-04	gb:NM_014413.2 /DEF=Homo sapiens heme-regulated initiation factor 2-alpha kinase (HRI), mRNA. /FEA=mRNA /GEN=HRI /PROD=heme-regulated initiation factor 2-alpha kinase /DB_XREF=gi:11125767 /UG=Hs.258730 heme-regulated initiation factor 2-alpha kinase /FL=gb:NM_014413.2 gb:AL136563.1 gb:AF147094.1 gb:AF255050.1 gb:AF116634.1 gb:AF183414.1	NM_014413		NP_055228
217737_x_at	1.26E-04	gb:NM_016407.1 /DEF=Homo sapiens hypothetical protein (HSPC164), mRNA. /FEA=mRNA /GEN=HSPC164 /PROD=hypothetical protein /DB_XREF=gi:7705482 /UG=Hs.182281 hypothetical protein /FL=gb:BC003359.1 gb:AF212244.1 gb:AF117231.1 gb:AF161513.1 gb:AF161518.1 gb:NM_016407.1	NM_016407		NP_057491
217739_s_at	0.033381	gb:NM_005746.1 /DEF=Homo sapiens pre-B-cell colony-enhancing factor (PBEF), mRNA. /FEA=mRNA /GEN=PBEF /PROD=pre-B-cell colony-enhancing factor /DB_XREF=gi:5031976 /UG=Hs.239138 pre-B-cell colony-enhancing factor /FL=gb:U02020.1 gb:NM_005746.1	NM_005746		NP_005737
217741_s_at	3.26E-04	zinc finger protein 216	AW471220	Hs.3776	NP_005998

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217742_s_at	0.004341	gb:NM_016628.1 /DEF=Homo sapiens hypothetical protein (LOC51322), mRNA. /FEA=mRNA /GEN=LOC51322 /PROD=hypothetical protein /DB_XREF=gi:7706169 /UG=Hs.70333 hypothetical protein /FL=gb:BC004258.1 gb:AF208858.1 gb:NM_016628.1	NM_016628		NP_567823
217743_s_at	1.26E-04	gb:NM_018247.1 /DEF=Homo sapiens hypothetical protein FLJ10856 (FLJ10856), mRNA. /FEA=mRNA /GEN=FLJ10856 /PROD=hypothetical protein FLJ10856 /DB_XREF=gi:8922719 /UG=Hs.108530 hypothetical protein FLJ10856 /FL=gb:NM_018247.1	NM_018247		NP_060717
217745_s_at	0.001116	gb:NM_025146.1 /DEF=Homo sapiens hypothetical protein FLJ13194 (FLJ13194), mRNA. /FEA=mRNA /GEN=FLJ13194 /PROD=hypothetical protein FLJ13194 /DB_XREF=gi:13376734 /UG=Hs.288932 hypothetical protein FLJ13194 /FL=gb:NM_025146.1	NM_025146		NP_079422
217746_s_at	1.26E-04	gb:NM_013374.1 /DEF=Homo sapiens programmed cell death 6 interacting protein (PDCD6IP), mRNA. /FEA=mRNA /GEN=PDCD6IP /PROD=programmed cell death 6-interacting protein /DB_XREF=gi:7019486 /UG=Hs.9663 programmed cell death 6-interacting protein /FL=gb:AF349951.1 gb:AF151793.1 gb:NM_013374.1	NM_013374		NP_037506
217747_s_at	0.041055	gb:NM_001013.1 /DEF=Homo sapiens ribosomal protein S9 (RPS9), mRNA. /FEA=mRNA /GEN=RPS9 /PROD=ribosomal protein S9 /DB_XREF=gi:4506744 /UG=Hs.180920 ribosomal protein S9 /FL=gb:BC000802.1 gb:NM_001013.1 gb:U14971.1	NM_001013		NP_001004

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217748_at	1.26E-04	gb:NM_015999.1 /DEF=Homo sapiens CGI-45 protein (LOC51094), mRNA. /FEA=mRNA /GEN=LOC51094 /PROD=CGI-45 protein /DB_XREF=gi:7705760 /UG=Hs.5298 CGI-45 protein /FL=gb:AF151803.1 gb:NM_015999.1	NM_015999		NP_057083
217755_at	1.26E-04	gb:NM_016185.1 /DEF=Homo sapiens hematological and neurological expressed 1 (HN1), mRNA. /FEA=mRNA /GEN=HN1 /PROD=hematological and neurological expressed 1 /DB_XREF=gi:7705876 /UG=Hs.109706 hematological and neurological expressed 1. /FL=gb:AF060925.1 gb:BC001420.1 gb:AF177862.1 gb:NM_016185.1	NM_016185		NP_057269
217758_s_at	0.00161	gb:NM_020123.1 /DEF=Homo sapiens endomembrane protein emp70 precursor isolog (LOC56889), mRNA. /FEA=mRNA /GEN=LOC56889 /PROD=endomembrane protein emp70 precursor isolog /DB_XREF=gi:10047129 /UG=Hs.8203 endomembrane protein emp70 precursor isolog /FL=gb:NM_020123.1 gb:AF160213.1 gb:AF269150.1	NM_020123		NP_064508
217762_s_at	1.26E-04	RAB31, member RAS oncogene family	BE789881	Hs.223025	NP_006859
217764_s_at	1.26E-04	gb:AF183421.1 /DEF=Homo sapiens small GTP-binding protein rab22b mRNA, complete cds. /FEA=mRNA /PROD=small GTP-binding protein rab22b /DB_XREF=gi:9963780 /UG=Hs.223025 RAB31, member RAS oncogene family /FL=gb:AF234995.1 gb:BC001148.1 gb:U59877.1 gb:U57091.1 gb:NM_006868.1 gb:AF183421.1	AF183421		NP_006859

Gene Identifier	p-value	D scription	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217766_s_at	0.033381	gb:NM_014313.1 /DEF=Homo sapiens small membrane protein 1 (SMP1), mRNA. /FEA=mRNA /GEN=SMP1 /PROD=small membrane protein 1 /DB_XREF=gi:7657594 /UG=Hs.107979 small membrane protein 1 /FL=gb:AL136627.1 gb:AF081282.1 gb:NM_014313.1	NM_014313		NP_055128
217768_at	0.026891	gb:NM_016039.1 /DEF=Homo sapiens CGI-99 protein (LOC51637), mRNA. /FEA=mRNA /GEN=LOC51637. /PROD=CGI-99 protein /DB_XREF=gi:7706321 /UG=Hs.110803 CGI-99 protein /FL=gb:BC001722.1 gb:AF151857.1 gb:AF100755.1 gb:NM_016039.1	NM_016039		NP_057123
217769_s_at	0.004341	gb:NM_015932.1 /DEF=Homo sapiens hypothetical protein (HSPC014), mRNA. /FEA=mRNA /GEN=HSPC014 /PROD=hypothetical protein /DB_XREF=gi:7705428 /UG=Hs.279813 hypothetical protein /FL=gb:AF275807.1 gb:BC003390.1 gb:AF077200.1 gb:AF125097.1 gb:NM_015932.1	NM_015932		NP_057016
217772_s_at	0.004733	gb:NM_014342.1 /DEF=Homo sapiens mitochondrial carrier homolog 2 (MTCH2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=MTCH2 /PROD=mitochondrial carrier homolog 2 /DB_XREF=gi:7657346 /UG=Hs.279609 mitochondrial carrier homolog 2 /FL=gb:BC000875.1 gb:AF085361.1 gb:AF176008.1 gb:NM_014342.1	NM_014342		NP_055157

Gen Id ntifi r	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217774_s_at	0.026891	gb:NM_016404.1 /DEF=Homo sapiens hypothetical protein (HSPC152), mRNA. /FEA=mRNA /GEN=HSPC152 /PROD=hypothetical protein /DB_XREF=gi:7705476 /UG=Hs.79259 hypothetical protein /FL=gb:AF110774.1 gb:AF161501.1 gb:NM_016404.1 gb:AF229068.1	NM_016404		NP_057488
217780_at	0.041055	gb:NM_016145.1 /DEF=Homo sapiens PTD008 protein (PTD008), mRNA. /FEA=mRNA /GEN=PTD008 /PROD=PTD008 protein /DB_XREF=gi:7706664 /UG=Hs.108969 PTD008 protein /FL=gb:AF059620.1 gb:AF151898.1 gb:AF078861.1 gb:NM_016145.1	NM_016145		NP_057229
217781_s_at	1.26E-04	gb:NM_022473.1 /DEF=Homo sapiens zinc finger protein, 106 (ZFP106), mRNA. /FEA=mRNA /GEN=ZFP106 /PROD=zinc finger protein 106 /DB_XREF=gi:11968022 /UG=Hs.15220 zinc finger protein 106 /FL=gb:AF205632.1 gb:NM_022473.1	NM_022473		NP_071918
217783_s_at	0.013223	gb:NM_016061.1 /DEF=Homo sapiens CGI-127 protein (LOC51646), mRNA. /FEA=mRNA /GEN=LOC51646 /PROD=CGI-127 protein /DB_XREF=gi:7706340 /UG=Hs.184542 CGI-127 protein /FL=gb:BC000836.1 gb:AF151885.1 gb:NM_016061.1	NM_016061		NP_057145
217786_at	9.79E-04	gb:NM_006109.1 /DEF=Homo sapiens skb1 (S. pombe) homolog (SKB1), mRNA. /FEA=mRNA /GEN=SKB1 /PROD=skb1 (S. pombe) homolog /DB_XREF=gi:5174682 /UG=Hs.12912 skb1 (S. pombe) homolog /FL=gb:AF015913.1 gb:NM_006109.1 gb:AF167572.1	NM_006109		NP_006100

Gen Id ntifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
217788_s_at	7.58E-04	gb:NM_004481.2 /DEF=Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 (GalNAc-T2) (GALNT2), mRNA. /FEA=mRNA /GEN=GALNT2 /PROD=polypeptide N-acetylgalactosaminyltransferase 2 /DB_XREF=gi:9945385 /UG=Hs.130181 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 (GalNAc-T2) /FL=gb:NM_004481.2	NM_004481		NP_004472
217794_at	7.58E-04	gb:NM_018457.1 /DEF=Homo sapiens DKFZp564J157 protein (DKFZP564J157), mRNA. /FEA=mRNA /GEN=DKFZP564J157 /PROD=DKFZp564J157 protein /DB_XREF=gi:8922156 /UG=Hs.63042 DKFZp564J157 protein /FL=gb:AF217517.1 gb:NM_018457.1	NM_018457		NP_060927
217795_s_at	0.005852	hypothetical protein MGC3222	W74580	Hs.323193	NP_077310
217797_at	0.007779	gb:NM_016406.1 /DEF=Homo sapiens hypothetical protein (HSPC155), mRNA. /FEA=mRNA /GEN=HSPC155 /PROD=hypothetical protein /DB_XREF=gi:7705480 /UG=Hs.177507 hypothetical protein /FL=gb:BC005187.1 gb:AF151884.1 gb:AF161504.1 gb:NM_016406.1	NM_016406		NP_057490
217799_x_at	0.027239	gb:NM_003344.1 /DEF=Homo sapiens ubiquitin-conjugating enzyme E2H (homologous to yeast UBC8) (UBE2H), mRNA. /FEA=mRNA /GEN=UBE2H /PROD=ubiquitin-conjugating enzyme E2H (homologous to yeast UBC8) /DB_XREF=gi:4507782 /UG=Hs.28505 ubiquitin-conjugating enzyme E2H (homologous to yeast UBC8) /FL=gb:NM_003344.1	NM_003344		NP_003335

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217800_s_at	0.001116	gb:NM_030571.1 /DEF=Homo sapiens hypothetical protein MGC10924 similar to Nedd4 WW-binding protein 5 (MGC10924), mRNA. /FEA=mRNA /GEN=MGC10924 /PROD=hypothetical protein MGC10924 similar to Nedd4WW-binding protein 5 /DB_XREF=gi:13386479 /UG=Hs.9788 hypothetical protein MGC10924 similar to Nedd4 WW-binding protein 5 /FL=gb:BC004317.1 gb:NM_030571.1	NM_030571		NP_085048
217801_at	0.003171	gb:NM_006886.1 /DEF=Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F1 complex, epsilon subunit (ATP5E), mRNA. /FEA=mRNA /GEN=ATP5E /PROD=ATP synthase, H ⁺ transporting, mitochondrial F1 complex, epsilon subunit /DB_XREF=gi:5901895 /UG=Hs.177530 ATP synthase, H ⁺ transporting, mitochondrial F1 complex, epsilon subunit /FL=gb:BC001690.1 gb:BC003671.1 gb:AF077045.1 gb:NM_006886.1 gb:AF052955.1	NM_006886		NP_008817
217802_s_at	1.26E-04	gb:NM_022731.1 /DEF=Homo sapiens similar to rat nuclear ubiquitous casein kinase 2 (NUCKS), mRNA. /FEA=mRNA /GEN=NUCKS /PROD=similar to rat nuclear ubiquitous casein kinase2 /DB_XREF=gi:12232386 /UG=Hs.118064 similar to rat nuclear ubiquitous casein kinase 2 /FL=gb:NM_022731.1 gb:BC000805.1	NM_022731		NP_073568
217803_at	1.26E-04	gb:NM_022130.1 /DEF=Homo sapiens Golgi protein (GPP34), mRNA. /FEA=mRNA /GEN=GPP34 /PROD=Golgi protein /DB_XREF=gi:11545858 /UG=Hs.18271 Golgi protein /FL=gb:NM_022130.1	NM_022130		NP_071413

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217805_at	0.003171	gb:NM_004516.1 /DEF=Homo sapiens interleukin enhancer binding factor 3, 90kD (ILF3), mRNA. /FEA=mRNA /GEN=ILF3 /PROD=interleukin enhancer binding factor 3, 90kD /DB_XREF=gi:4758603 /UG=Hs.256583 interleukin enhancer binding factor 3, 90kD /FL=gb:BC003086.1 gb:NM_004516.1 gb:U10324.1 gb:AF167570.1 gb:NM_012218.1	NM_004516		NP_703194
217807_s_at	0.016934	gb:NM_015710.1 /DEF=Homo sapiens glioma tumor suppressor candidate region gene 2 (GLTSCR2), mRNA. /FEA=mRNA /GEN=GLTSCR2 /PROD=glioma tumor suppressor candidate region gene 2 /DB_XREF=gi:7657129 /UG=Hs.2237 glioma tumor suppressor candidate region gene 2 /FL=gb:AF182076.1 gb:NM_015710.1	NM_015710		NP_056525
217811_at	1.26E-04	gb:NM_016275.1 /DEF=Homo sapiens selenoprotein T (LOC51714), mRNA. /FEA=mRNA /GEN=LOC51714 /PROD=selenoprotein T /DB_XREF=gi:7706470 /UG=Hs.8148 selenoprotein T /FL=gb:AF131856.1 gb:AF195141.1 gb:NM_016275.1	NM_016275		NP_057359
217812_at	0.016934	gb:NM_016258.1 /DEF=Homo sapiens high-glucose-regulated protein 8 (HGRG8), mRNA. /FEA=mRNA /GEN=HGRG8 /PROD=high-glucose-regulated protein 8 /DB_XREF=gi:7705410 /UG=Hs.20993 high-glucose-regulated protein 8 /FL=gb:BC002559.1 gb:AF155095.1 gb:AF192968.1 gb:NM_016258.1	NM_016258		NP_057342

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
217814_at	1.26E-04	gb:NM_020198.1 /DEF=Homo sapiens GK001 protein (GK001), mRNA. /FEA=mRNA /GEN=GK001 /PROD=GK001 protein /DB_XREF=gi:9910241 /UG=Hs.8207 GK001 protein /FL=gb:AF113221.1 gb:BC001300.1 gb:AF226054.1 gb:NM_020198.1	NM_020198		NP_064583
217815_at	0.00161	gb:NM_007192.1 /DEF=Homo sapiens chromatin-specific transcription elongation factor, 140 kDa subunit (FACTP140), mRNA. /FEA=mRNA /GEN=FACTP140 /PROD=chromatin-specific transcription elongation factor, 140 kDa subunit. /DB_XREF=gi:6005756 /UG=Hs.14963 chromatin-specific transcription elongation factor, 140 kDa subunit /FL=gb:AF152961.1 gb:NM_007192.1	NM_007192		NP_009123
217819_at	1.26E-04	gb:NM_016099.1 /DEF=Homo sapiens HSPC041 protein (LOC51125), mRNA. /FEA=mRNA /GEN=LOC51125 /PROD=HSPC041 protein /DB_XREF=gi:7705820 /UG=Hs.7953 HSPC041 protein /FL=gb:AF125102.1 gb:NM_016099.1	NM_016099		NP_057183
217823_s_at	1.26E-04	Consensus includes gb:AL562528 /FEA=EST /DB_XREF=gi:12911037 /DB_XREF=est:AL562528 /CLONE=CS0DC012YI08 (3 prime) /UG=Hs.184325 CGI-76 protein /FL=gb:AF151834.1 gb:AF161502.1 gb:AF151039.1 gb:NM_016021.1	NM_016021		NP_057420

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
217824_at	0.02145	Consensus includes gb:AW500009 /FEA=EST /DB_XREF=gi:7112213 /DB_XREF=est:UI-HF-BN0-aki-e- 09-0-UI.r1 /CLONE=IMAGE:3077105 /UG=Hs.184325 CGI-76 protein /FL=gb:AF151834.1 gb:AF161502.1 gb:AF151039.1 gb:NM_016021.1	NM_016021		NP_057420
217825_s_at	0.00228	gb:AF151039.1 /DEF=Homo sapiens HSPC205 mRNA; complete cds. /FEA=mRNA /PROD=HSPC205 /DB_XREF=gi:7106799 /UG=Hs.184325 CGI-76 protein /FL=gb:AF151834.1 gb:AF161502.1 gb:AF151039.1 gb:NM_016021.1	AF151039		NP_057420
217826_s_at	7.58E-04	gb:NM_016021.1 /DEF=Homo sapiens CGI-76 protein (LOC51632), mRNA. /FEA=mRNA /GEN=LOC51632 /PROD=CGI-76 protein /DB_XREF=gi:7706311 /UG=Hs.184325 CGI-76 protein /FL=gb:AF151834.1 gb:AF161502.1 gb:AF151039.1 gb:NM_016021.1	NM_016021		NP_057420
217827_s_at	2.05E-04	gb:NM_016630.2 /DEF=Homo sapiens acid cluster protein 33 (ACP33), mRNA. /FEA=mRNA /GEN=ACP33 /PROD=acid cluster protein 33 /DB_XREF=gi:13699804 /UG=Hs.71475 acid cluster protein 33 /FL=gb:BC000244.1 gb:NM_016630.2 gb:AF208861.1	NM_016630		NP_057714
217835_x_at	3.26E-04	gb:NM_018840.1 /DEF=Homo sapiens putative Rab5-interacting protein (LOC55969), mRNA. /FEA=mRNA /GEN=LOC55969 /PROD=putative Rab5-interacting protein /DB_XREF=gi:10047115 /UG=Hs.184062 putative Rab5- interacting protein /FL=gb:NM_018840.1 gb:AF274936.1 gb:AF112213.1	NM_018840		NP_061328

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
217836_s_at	0.005852	gb:NM_018253.1 /DEF=Homo sapiens hypothetical protein FLJ10875 (FLJ10875), mRNA. /FEA=mRNA /GEN=FLJ10875 /PROD=hypothetical protein FLJ10875 /DB_XREF=gi:8922730 /UG=Hs.18851 hypothetical protein FLJ10875 /FL=gb:BC001655.1 gb:BC001843.1 gb:NM_018253.1	NM_018253		NP_620832
217837_s_at	1.26E-04	gb:NM_016079.1 /DEF=Homo sapiens CGI-149 protein (LOC51652), mRNA. /FEA=mRNA /GEN=LOC51652 /PROD=CGI-149 protein /DB_XREF=gi:7706352 /UG=Hs.189658 CGI-149 protein /FL=gb:BC004419.1 gb:AF151907.1 gb:AF219226.1 gb:NM_016079.1	NM_016079		NP_057163
217838_s_at	0.033381	gb:NM_016337.1 /DEF=Homo sapiens RNB6 (RNB6), mRNA. /FEA=mRNA /GEN=RNB6 /PROD=RNB6 /DB_XREF=gi:7706686 /UG=Hs.241471 RNB6 /FL=gb:AF052504.1 gb:NM_016337.1	NM_016337		NP_057421
217839_at	6.42E-04	gb:NM_006070.1 /DEF=Homo sapiens TRK-fused gene (TFG), mRNA. /FEA=mRNA /GEN=TFG /PROD=TRK-fused gene /DB_XREF=gi:5174718 /UG=Hs.250897 TRK-fused gene /FL=gb:NM_006070.1	NM_006070		NP_006061
217843_s_at	3.26E-04	gb:NM_014166.1 /DEF=Homo sapiens HSPC126 protein (HSPC126), mRNA. /FEA=mRNA /GEN=HSPC126 /PROD=HSPC126 protein /DB_XREF=gi:7661787 /UG=Hs.181112 HSPC126 protein /FL=gb:AF300618.1 gb:BC005189.1 gb:AF161475.1 gb:AF230381.1 gb:NM_014166.1	NM_014166		NP_054885

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217844_at	0.003529	gb:NM_021198.1 /DEF=Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA. /FEA=mRNA /GEN=NLI-IF /PROD=nuclear LIM interactor-interacting factor /DB_XREF=gi:10864008 /UG=Hs.283724 nuclear LIM interactor-interacting factor /FL=gb:AF229162.1 gb:NM_021198.1	NM_021198		NP_067021
217845_x_at	0.00161	gb:NM_014056.1 /DEF=Homo sapiens DKFZP564K247 protein (DKFZP564K247), mRNA. /FEA=mRNA /GEN=DKFZP564K247 /PROD=DKFZP564K247 protein /DB_XREF=gi:7661619. /UG=Hs.7917 DKFZP564K247 protein /FL=gb:BC000601.1 gb:AF077034.1 gb:AF145385.1 gb:AL110233.1 gb:NM_014056.1	NM_014056		NP_054775
217852_s_at	1.26E-04	gb:NM_018184.1 /DEF=Homo sapiens hypothetical protein FLJ10702 (FLJ10702), mRNA. /FEA=mRNA /GEN=FLJ10702 /PROD=hypothetical protein FLJ10702 /DB_XREF=gi:8922600 /UG=Hs.104222 hypothetical protein FLJ10702 /FL=gb:NM_018184.1	NM_018184		NP_060654
217854_s_at	0.022795	gb:NM_002695.1 /DEF=Homo sapiens polymerase (RNA) II (DNA directed) polypeptide E (25kD) (POLR2E), mRNA. /FEA=mRNA /GEN=POLR2E /PROD=polymerase (RNA) II (DNA directed) polypeptide E(25kD) /DB_XREF=gi:4505944 /UG=Hs.24301 polymerase (RNA) II (DNA directed) polypeptide E (25kD) /FL=gb:BC004441.1 gb:NM_002695.1 gb:J04965.1 gb:D38251.1	NM_002695		NP_002686

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
217855_x_at	0.004341	gb:NM_016547.1 /DEF=Homo sapiens calcium binding protein Cab45 precursor, (LOC51767), mRNA. /FEA=mRNA /GEN=LOC51767 /PROD=calcium binding protein Cab45 precursor, /DB_XREF=gi:7706572 /UG=Hs.42806 calcium binding protein Cab45 precursor, /FL=gb:AF178986.1 gb:NM_016547.1	NM_016547		NP_057631
217862_at	3.26E-04	Consensus includes gb:N24868 /FEA=EST /DB_XREF=gi:1139018 /DB_XREF=est:yx70h06.s1 /CLONE=IMAGE:267131 /UG=Hs.75251 DEADH (Asp-Glu-Ala-AspHis) box binding protein 1 /FL=gb:AF077951.1 gb:AF167160.1 gb:NM_016166.1	NM_016166		NP_057250
217863_at	0.00228	Consensus includes gb:AI348378 /FEA=EST /DB_XREF=gi:4085584 /DB_XREF=est:qo20h06.x1 /CLONE=IMAGE:1909115 /UG=Hs.75251 DEADH (Asp-Glu-Ala-AspHis) box binding protein 1 /FL=gb:AF077951.1 gb:AF167160.1 gb:NM_016166.1	NM_016166		NP_057250
217865_at	2.05E-04	gb:NM_018434.1 /DEF=Homo sapiens goliath protein (LOC55819), mRNA. /FEA=mRNA /GEN=LOC55819 /PROD=goliath protein /DB_XREF=gi:10092650 /UG=Hs.102737 goliath protein /FL=gb:NM_018434.1	NM_018434		NP_060904
217868_s_at	1.26E-04	gb:NM_016025.1 /DEF=Homo sapiens CGI-81 protein (DREV1), mRNA. /FEA=mRNA /GEN=DREV1 /PROD=CGI-81 protein /DB_XREF=gi:7705788 /UG=Hs.279583 CGI-81 protein /FL=gb:BC000195.1 gb:AF151839.1 gb:NM_016025.1	NM_016025		NP_057109

Gen Id ntifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217869_at	1.26E-04	gb:NM_016142.1 /DEF=Homo sapiens steroid dehydrogenase homolog (LOC51144), mRNA. /FEA=mRNA /GEN=LOC51144 /PROD=steroid dehydrogenase homolog /DB_XREF=gi:7705854 /UG=Hs.279617 steroid dehydrogenase homolog /FL=gb:AF078850.1 gb:NM_016142.1	NM_016142		NP_057226
217873_at	1.26E-04	gb:NM_016289.1 /DEF=Homo sapiens MO25 protein (LOC51719), mRNA. /FEA=mRNA /GEN=LOC51719 /PROD=MO25 protein /DB_XREF=gi:7706480 /UG=Hs.6406 MO25 protein /FL=gb:AF113536.1 gb:NM_016289.1	NM_016289		NP_057373
217874_at	1.26E-04	gb:NM_003849.1 /DEF=Homo sapiens succinate-CoA ligase, GDP-forming, alpha subunit (SUCLG1), mRNA. /FEA=mRNA /GEN=SUCLG1 /PROD=succinate-CoA ligase, GDP-forming, alpha subunit /DB_XREF=gi:11321580 /UG=Hs.7043 succinate-CoA ligase, GDP-forming, alpha subunit /FL=gb:NM_003849.1 gb:BC000504.1 gb:AF104921.2	NM_003849		NP_003840
217877_s_at	0.02145	gb:NM_021639.1 /DEF=Homo sapiens hypothetical protein SP192 (SP192), mRNA. /FEA=mRNA /GEN=SP192 /PROD=hypothetical protein SP192 /DB_XREF=gi:11056015 /UG=Hs.169854 hypothetical protein SP192 /FL=gb:NM_021639.1	NM_021639		NP_067652
217880_at	0.005852	cell division cycle 27	AI203880	Hs.408978	NP_001247
217882_at	0.00161	gb:NM_018447.1 /DEF=Homo sapiens 30 kDa protein (LOC55831), mRNA. /FEA=mRNA /GEN=LOC55831 /PROD=30 kDa protein /DB_XREF=gi:8923856 /UG=Hs.283714 30 kDa protein /FL=gb:AF157321.1 gb:NM_018447.1	NM_018447		NP_060917

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217883_at	1.26E-04	gb:NM_015702.1 /DEF=Homo sapiens hypothetical protein (CL25022), mRNA. /FEA=mRNA /GEN=CL25022 /PROD=hypothetical protein /DB_XREF=gi:7661547 /UG=Hs.5324 hypothetical protein /FL=gb:AF060224.1 gb:BC000932.2 gb:AF131802.1 gb:AF161510.1 gb:NM_015702.1	NM_015702		NP_056517
217885_at	3.26E-04	gb:NM_018085.1 /DEF=Homo sapiens hypothetical protein FLJ10402 (FLJ10402), mRNA. /FEA=mRNA /GEN=FLJ10402 /PROD=hypothetical protein FLJ10402 /DB_XREF=gi:8922404 /UG=Hs.7579 hypothetical protein FLJ10402 /FL=gb:BC003604.1 gb:NM_018085.1	NM_018085		NP_060555
217886_at	0.001116	epidermal growth factor receptor pathway substrate 15	BF213575	Hs.389343	NP_001972
217887_s_at	0.007779	gb:NM_001981.1 /DEF=Homo sapiens epidermal growth factor receptor pathway substrate 15 (EPS15), mRNA. /FEA=mRNA /GEN=EPS15 /PROD=epidermal growth factor receptor pathway substrate 15 /DB_XREF=gi:4503592 /UG=Hs.79095 epidermal growth factor receptor pathway substrate 15 /FL=gb:NM_001981.1 gb:U07707.1	NM_001981		NP_001972
217898_at	2.05E-04	gb:NM_020154.1 /DEF=Homo sapiens chromosome 11 hypothetical protein ORF3 (LOC56851), mRNA. /FEA=mRNA /GEN=LOC56851 /PROD=chromosome 11 hypothetical protein ORF3 /DB_XREF=gi:9910345 /UG=Hs.4245 chromosome 11 hypothetical protein ORF3 /FL=gb:AF242729.1 gb:NM_020154.1	NM_020154		NP_064539

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
217900_at	0.005852	gb:NM_018060.1 /DEF=Homo sapiens hypothetical protein FLJ10326 (FLJ10326), mRNA. /FEA=mRNA /GEN=FLJ10326 /PROD=hypothetical protein FLJ10326 /DB_XREF=gi:8922355 /UG=Hs.262823 hypothetical protein FLJ10326 /FL=gb:NM_018060.1	NM_018060		NP_060530
217903_at	0.016934	gb:NM_013403.1 /DEF=Homo sapiens zinedin (ZIN), mRNA. /FEA=mRNA /GEN=ZIN /PROD=zinedin /DB_XREF=gi:7019572 /UG=Hs.108665 zinedin /FL=gb:AF212940.1 gb:NM_013403.1	NM_013403		NP_037535
217905_at	7.58E-04	gb:NM_024834.1 /DEF=Homo sapiens hypothetical protein FLJ13081 (FLJ13081), mRNA. /FEA=mRNA /GEN=FLJ13081 /PROD=hypothetical protein FLJ13081 /DB_XREF=gi:13376242 /UG=Hs.180638 hypothetical protein FLJ13081 /FL=gb:BC004183.1 gb:NM_024834.1	NM_024834		NP_079110
217907_at	7.58E-04	gb:NM_014161.1 /DEF=Homo sapiens HSPC071 protein (HSPC071), mRNA. /FEA=mRNA /GEN=HSPC071 /PROD=HSPC071 protein /DB_XREF=gi:7661777 /UG=Hs.23038 HSPC071 protein /FL=gb:AL136633.1 gb:BC001623.1 gb:AF161556.1 gb:NM_014161.1	NM_014161		NP_054880
217908_s_at	1.26E-04	gb:NM_018442.1 /DEF=Homo sapiens PC326 protein (PC326), mRNA. /FEA=mRNA /GEN=PC326 /PROD=PC326 protein /DB_XREF=gi:8923955 /UG=Hs.279882 PC326 protein /FL=gb:AL136738.1 gb:AF150734.1 gb:NM_018442.1	NM_018442		NP_060912

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217910_x_at	1.26E-04	gb:NM_013383.1 /DEF=Homo sapiens MAX-like bHLHZIP protein (MLX), mRNA. /FEA=mRNA /GEN=MLX /PROD=MAX-like bHLHZIP protein /DB_XREF=gi:7019460 /UG=Hs.78185 MAX-like bHLHZIP protein /FL=gb:AF213668.1 gb:AF203978.1 gb:NM_013383.1	NM_013383		NP_733753
217911_s_at	0.003171	gb:NM_004281.1 /DEF=Homo sapiens BCL2-associated athanogene 3 (BAG3), mRNA. /FEA=mRNA /GEN=BAG3 /PROD=BCL2-associated athanogene 3 /DB_XREF=gi:6631072 /UG=Hs.15259 BCL2-associated athanogene 3 /FL=gb:AF095193.2 gb:NM_004281.1 gb:AF127139.1 gb:AF071218.2	NM_004281		NP_004272
217913_at	0.013223	gb:NM_013245.1 /DEF=Homo sapiens vacuolar sorting protein 4 (VPS4), mRNA. /FEA=mRNA /GEN=VPS4 /PROD=vacuolar sorting protein 4 /DB_XREF=gi:7019568 /UG=Hs.234839 vacuolar sorting protein 4 /FL=gb:AF159063.1 gb:AF112215.1 gb:NM_013245.1	NM_013245		NP_037377
217915_s_at	0.016934	gb:NM_016304.1 /DEF=Homo sapiens 60S ribosomal protein L30 isolog (LOC51187), mRNA. /FEA=mRNA /GEN=LOC51187 /PROD=60S ribosomal protein L30 isolog /DB_XREF=gi:10047101 /UG=Hs.284162 60S ribosomal protein L30 isolog /FL=gb:NM_016304.1 gb:AF060926.1 gb:AF212226.1 gb:BC005344.1 gb:AF201949.1 gb:AF165521.1	NM_016304		NP_057388

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217916_s_at	0.00161	gb:NM_016623.1 /DEF=Homo sapiens hypothetical protein (BM-009), mRNA. /FEA=mRNA /GEN=BM-009 /PROD=hypothetical protein /DB_XREF=gi:7705303 /UG=Hs.92918 hypothetical protein /FL=gb:BC003599.1 gb:AF208851.1 gb:NM_016623.1	NM_016623		NP_057707
217917_s_at	5.03E-04	gb:NM_014183.1 /DEF=Homo sapiens HSPC162 protein (HSPC162), mRNA. /FEA=mRNA /GEN=HSPC162 /PROD=HSPC162 protein /DB_XREF=gi:7661821 /UG=Hs.100002 HSPC162 protein /FL=gb:BC002481.1 gb:AY026513.1 gb:AF161511.1 gb:NM_014183.1 gb:AF165516.1	NM_014183		NP_808853
217918_at	1.26E-04	gb:NM_014183.1 /DEF=Homo sapiens HSPC162 protein (HSPC162), mRNA. /FEA=mRNA /GEN=HSPC162 /PROD=HSPC162 protein /DB_XREF=gi:7661821 /UG=Hs.100002 HSPC162 protein /FL=gb:BC002481.1 gb:AY026513.1 gb:AF161511.1 gb:NM_014183.1 gb:AF165516.1	NM_014183		NP_808853
217923_at	0.02145	gb:NM_012392.1 /DEF=Homo sapiens PEF protein with a long N-terminal hydrophobic domain (peflin) (PEF), mRNA. /FEA=mRNA /GEN=PEF /PROD=peflin /DB_XREF=gi:6912581 /UG=Hs.241531 peflin /FL=gb:BC002773.1 gb:AB018357.1 gb:AB026628.1 gb:NM_012392.1	NM_012392		NP_036524
217925_s_at	0.022752	gb:NM_022758.1 /DEF=Homo sapiens hypothetical protein FLJ22195 (FLJ22195), mRNA. /FEA=mRNA /GEN=FLJ22195 /PROD=hypothetical protein FLJ22195 /DB_XREF=gi:12232426 /UG=Hs.25999 hypothetical protein FLJ22195 /FL=gb:NM_022758.1	NM_022758		NP_073595

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217927_at	3.26E-04	gb:NM_014041.1 /DEF=Homo sapiens HSPC033 protein (HSPC033), mRNA. /FEA=mRNA /GEN=HSPC033 /PROD=HSPC033 protein /DB_XREF=gi:7661745 /UG=Hs.11125 HSPC033 protein /FL=gb:BC000884.1 gb:AF092138.1 gb:NM_014041.1	NM_014041		NP_054760
217928_s_at	0.041055	gb:NM_018312.2 /DEF=Homo sapiens chromosome 11 open reading frame 23 (C11ORF23), mRNA. /FEA=mRNA /GEN=C11ORF23 /PROD=sporulation-induced transcript 4-associated protein /DB_XREF=gi:13489082 /UG=Hs.180817 chromosome 11 open reading frame 23 /FL=gb:AF264779.1 gb:NM_018312.2	NM_018312		NP_060782
217930_s_at	0.009689	gb:NM_019009.1 /DEF=Homo sapiens TOLLIP protein (LOC54472), mRNA. /FEA=mRNA /GEN=LOC54472 /PROD=TOLLIP protein /DB_XREF=gi:9506854 /UG=Hs.25413 TOLLIP protein /FL=gb:AL136835.1 gb:BC004420.1 gb:NM_019009.1	NM_019009		NP_061882
217936_at	0.003171	Rho GTPase activating protein 5	AW044631	Hs.267831	
217938_s_at	0.00228	gb:NM_020122.1 /DEF=Homo sapiens potassium channel modulatory factor (DKFZP434L1021), mRNA. /FEA=mRNA /GEN=DKFZP434L1021 /PROD=potassium channel modulatory factor /DB_XREF=gi:10047127 /UG=Hs.5392 potassium channel modulatory factor /FL=gb:NM_020122.1 gb:BC000178.2	NM_020122		NP_064507

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217939_s_at	7.58E-04	gb:NM_017657.1 /DEF=Homo sapiens hypothetical protein FLJ20080 (FLJ20080), mRNA. /FEA=mRNA /GEN=FLJ20080 /PROD=hypothetical protein FLJ20080 /DB_XREF=gi:8923087 /UG=Hs.7942 hypothetical protein FLJ20080 /FL=gb:NM_017657.1	NM_017657		NP_060127
217941_s_at	0.00228	gb:NM_018695.1 /DEF=Homo sapiens erbb2-interacting protein ERBIN (LOC55914), mRNA. /FEA=mRNA /GEN=LOC55914 /PROD=erbb2-interacting protein ERBIN /DB_XREF=gi:8923908 /UG=Hs.8117 erbb2-interacting protein ERBIN /FL=gb:AF263744.1 gb:NM_018695.1	NM_018695		NP_061165
217949_s_at	5.03E-04	gb:NM_024006.1 /DEF=Homo sapiens hypothetical protein IMAGE3455200 (IMAGE3455200), mRNA. /FEA=mRNA /GEN=IMAGE3455200 /PROD=hypothetical protein IMAGE3455200 /DB_XREF=gi:13124769 /UG=Hs.324844 hypothetical protein IMAGE3455200 /FL=gb:BC002911.1 gb:NM_024006.1	NM_024006		NP_076869
217951_s_at	5.03E-04	PHD finger protein 3	AW189430	Hs.78893	NP_055968
217952_x_at	0.016934	PHD finger protein 3	AW189430	Hs.78893	NP_055968
217954_s_at	0.026891	gb:NM_015153.1 /DEF=Homo sapiens KIAA0244 protein (KIAA0244), mRNA. /FEA=mRNA /GEN=KIAA0244 /PROD=KIAA0244 protein /DB_XREF=gi:7662017 /UG=Hs.78893 KIAA0244 protein /FL=gb:AF091622.1 gb:NM_015153.1	NM_015153		NP_055968

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217955_at	1.26E-04	gb:NM_015367.1 /DEF=Homo sapiens MIL1 protein (MIL1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=MIL1 /PROD=MIL1 protein /DB_XREF=gi:7662505 /UG=Hs.10267 MIL1 protein /FL=gb:AF325209.1 gb:AF146568.1 gb:NM_015367.1	NM_015367		NP_056182
217958_at	0.026891	gb:NM_016146.1 /DEF=Homo sapiens PTD009 protein (PTD009), mRNA. /FEA=mRNA /GEN=PTD009 /PROD=PTD009 protein /DB_XREF=gi:7706666 /UG=Hs.279901 PTD009 protein /FL=gb:AF151862.1 gb:AF078862.1 gb:AF161520.1 gb:NM_016146.1	NM_016146		NP_057230
217959_s_at	0.004341	gb:NM_016146.1 /DEF=Homo sapiens PTD009 protein (PTD009), mRNA. /FEA=mRNA /GEN=PTD009 /PROD=PTD009 protein /DB_XREF=gi:7706666 /UG=Hs.279901 PTD009 protein /FL=gb:AF151862.1 gb:AF078862.1 gb:AF161520.1 gb:NM_016146.1	NM_016146		NP_057230
217962_at	1.26E-04	gb:NM_018648.1 /DEF=Homo sapiens nucleolar protein family A, member 3 (HACA small nucleolar RNPs) (NOLA3), mRNA. /FEA=mRNA /GEN=NOLA3 /PROD=nucleolar protein family A, member 3 (HACA small nucleolar RNPs) /DB_XREF=gi:8923941 /UG=Hs.14317 nucleolar protein family A, member 3 (HACA small nucleolar RNPs) /FL=gb:AB043104.1 gb:NM_018648.1	NM_018648		NP_061118
217967_s_at	0.033381	gb:AF288391.1 /DEF=Homo sapiens C1orf24 mRNA, complete cds. /FEA=mRNA /PROD=C1orf24 /DB_XREF=gi:12620191 /UG=Hs.48778 niban protein /FL=gb:AB050477.1 gb:NM_022083.1 gb:AF288391.1	AF288391		NP_443198

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217969_at	0.041055	gb:NM_013265.2 /DEF=Homo sapiens chromosome 11 open reading frame2 (C11ORF2), mRNA. /FEA=mRNA /GEN=C11ORF2 /PROD=chromosome 11 open reading frame2 /DB_XREF=gi:8393008 /UG=Hs.5258 chromosome 11 open reading frame2 /FL=gb:AF024631.2 gb:NM_013265.2	NM_013265		NP_008917
217970_s_at	0.002313	gb:NM_015455.1 /DEF=Homo sapiens KIAA1194 protein (KIAA1194), mRNA. /FEA=mRNA /GEN=KIAA1194 /PROD=DKFZP564G1964 protein /DB_XREF=gi:7661613 /UG=Hs.5801 KIAA1194 protein /FL=gb:NM_015455.1	NM_015455		NP_056270
217971_at	1.26E-04	gb:NM_021970.1 /DEF=Homo sapiens MEK partner 1 (MP1), mRNA. /FEA=mRNA /GEN=MP1 /PROD=MEK partner 1 /DB_XREF=gi:11496276 /UG=Hs.6361 MEK partner 1 /FL=gb:AF130115.1 gb:NM_021970.1 gb:AF201947.1	NM_021970		NP_068805
217972_at	0.02145	gb:NM_017812.1 /DEF=Homo sapiens hypothetical protein FLJ20420 (FLJ20420), mRNA. /FEA=mRNA /GEN=FLJ20420 /PROD=hypothetical protein FLJ20420 /DB_XREF=gi:8923389 /UG=Hs.6693 hypothetical protein FLJ20420 /FL=gb:NM_017812.1	NM_017812		NP_060282
217977_at	0.00161	gb:NM_016332.1 /DEF=Homo sapiens selenoprotein X, 1 (SEPX1), mRNA. /FEA=mRNA /GEN=SEPX1 /PROD=selenoprotein X, 1 /DB_XREF=gi:7706510 /UG=Hs.279623 selenoprotein X, 1 /FL=gb:AF187272.1 gb:BC003127.1 gb:AF166124.1 gb:NM_016332.1	NM_016332		NP_057416

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217978_s_at	7.58E-04	gb:NM_017582.1 /DEF=Homo sapiens NICE-5 protein (HSA243666), mRNA. /FEA=mRNA /GEN=HSA243666 /PROD=NICE-5 protein. /DB_XREF=gi:8923743 /UG=Hs.284233 NICE-5 protein /FL=gb:AF116721.1 gb:NM_017582.1	NM_017582		NP_060052
217980_s_at	0.005852	gb:NM_017840.1 /DEF=Homo sapiens hypothetical protein FLJ20484 (FLJ20484), mRNA. /FEA=mRNA /GEN=FLJ20484 /PROD=hypothetical protein FLJ20484 /DB_XREF=gi:8923447 /UG=Hs.5080 hypothetical protein FLJ20484 /FL=gb:AF275806.1 gb:BC001040.1 gb:NM_017840.1 gb:AF183428.1	NM_017840		NP_060310
217981_s_at	0.004341	gb:NM_012192.1 /DEF=Homo sapiens fracture callus 1 (rat) homolog (FXC1), mRNA. /FEA=mRNA /GEN=FXC1 /PROD=fracture callus 1 (rat) homolog /DB_XREF=gi:6912381 /UG=Hs.54943 fracture callus 1 (rat) homolog /FL=gb:AF150105.1 gb:AF152355.1 gb:NM_012192.1 gb:AF183415.1	NM_012192		NP_036324
217983_s_at	0.007779	gb:NM_003730.2 /DEF=Homo sapiens ribonuclease 6 precursor (RNASE6PL), mRNA. /FEA=mRNA /GEN=RNASE6PL /PROD=ribonuclease 6 precursor /DB_XREF=gi:5231227 /UG=Hs.8297 ribonuclease 6 precursor /FL=gb:BC001660.1 gb:BC001819.1 gb:U85625.2 gb:NM_003730.2	NM_003730		NP_003721

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
217986_s_at	2.05E-04	gb:NM_013448.1 /DEF=Homo sapiens bromodomain adjacent to zinc finger domain, 1A (BAZ1A), mRNA. /FEA=mRNA. /GEN=BAZ1A /PROD=bromodomain adjacent to zinc finger domain, 1A /DB_XREF=gi:7304918 /UG=Hs.8858 bromodomain adjacent to zinc finger domain, 1A /FL=gb:AB032252.1 gb:NM_013448.1	NM_013448		NP_038476
217989_at	0.010205	gb:NM_016245.1 /DEF=Homo sapiens retinal short-chain dehydrogenasereductase retSDR2 (LOC51170), mRNA. /FEA=mRNA /GEN=LOC51170 /PROD=retinal short-chain dehydrogenasereductaseretSDR2 /DB_XREF=gi:7705904 /UG=Hs.12150 retinal short-chain dehydrogenasereductase. retSDR2 /FL=gb:AF126780.1 gb:NM_016245.1	NM_016245		NP_057329
217990_at	7.58E-04	gb:NM_016576.1 /DEF=Homo sapiens GMPR2 for guanosine monophosphate reductase isolog (LOC51292), mRNA. /FEA=mRNA /GEN=LOC51292 /PROD=GMPR2 for guanosine monophosphate reductaseisolog /DB_XREF=gi:7706108 /UG=Hs.234546 GMPR2 for guanosine monophosphate reductase isolog /FL=gb:AB032903.1 gb:NM_016576.1 gb:AF135159.1	NM_016576		NP_057660
217995_at	1.26E-04	gb:NM_021199.1 /DEF=Homo sapiens CGI-44 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA. /FEA=mRNA /GEN=CGI-44 /PROD=CGI-44 protein; sulfide dehydrogenase like(yeast) /DB_XREF=gi:10864010 /UG=Hs.8185 CGI-44 protein; sulfide dehydrogenase like (yeast) /FL=gb:NM_021199.1 gb:AF151802.1 gb:AF118085.1	NM_021199		NP_067022

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
218004_at	0.016934	gb:NM_018045.1 /DEF=Homo sapiens hypothetical protein FLJ10276 (FLJ10276), mRNA. /FEA=mRNA /GEN=FLJ10276 /PROD=hypothetical protein FLJ10276 /DB_XREF=gi:8922323 /UG=Hs.6937 hypothetical protein FLJ10276 /FL=gb:NM_018045.1	NM_018045		NP_060515
218006_s_at	0.001618	gb:NM_006963.1 /DEF=Homo sapiens zinc finger protein 22 (KOX 15) (ZNF22), mRNA. /FEA=mRNA /GEN=ZNF22 /PROD=zinc finger protein 22 (KOX 15) /DB_XREF=gi:5902159 /UG=Hs.108642 zinc finger protein 22 (KOX 15) /FL=gb:NM_006963.1	NM_006963		NP_008894
218008_at	7.58E-04	gb:NM_017994.1 /DEF=Homo sapiens hypothetical protein FLJ10099 (FLJ10099), mRNA. /FEA=mRNA /GEN=FLJ10099 /PROD=hypothetical protein FLJ10099 /DB_XREF=gi:8922228 /UG=Hs.278619 hypothetical protein FLJ10099 /FL=gb:NM_017994.1	NM_017994		NP_060464
218011_at	1.26E-04	gb:NM_024292.1 /DEF=Homo sapiens ubiquitin-like 5 (UBL5), mRNA. /FEA=mRNA /GEN=UBL5 /PROD=ubiquitin-like 5 /DB_XREF=gi:13236509 /UG=Hs.109701 ubiquitin-like 5 /FL=gb:AF313915.1 gb:NM_024292.1	NM_024292		NP_077268
218013_x_at	0.034221	gb:NM_016221.1 /DEF=Homo sapiens dynactin p62 subunit (LOC51164), mRNA. /FEA=mRNA /GEN=LOC51164 /PROD=dynactin p62 subunit /DB_XREF=gi:7705892 /UG=Hs.180952 dynactin 4 (p62) /FL=gb:AF195120.1 gb:NM_016221.1	NM_016221		NP_057305

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218017_s_at	0.026891	gb:NM_025070.1 /DEF=Homo sapiens hypothetical protein FLJ22242 (FLJ22242), mRNA. /FEA=mRNA /GEN=FLJ22242 /PROD=hypothetical protein FLJ22242 /DB_XREF=gi:13376612 /UG=Hs.288057 hypothetical protein FLJ22242 /FL=gb:NM_025070.1	NM_025070		NP_079346
218018_at	0.021448	Consensus includes gb:AW449022 /FEA=EST /DB_XREF=gi:6989798 /DB_XREF=est:UI-H-BI3-ake-f-11-0 UI.s1 /CLONE=IMAGE:2734220 /UG=Hs.4746 hypothetical protein FLJ21324 /FL=gb:NM_021941.1 gb:BC003651.1	NM_021941		NP_068760
218019_s_at	0.013223	gb:NM_021941.1 /DEF=Homo sapiens hypothetical protein FLJ21324 (FLJ21324), mRNA. /FEA=mRNA /GEN=FLJ21324 /PROD=hypothetical protein FLJ21324 /DB_XREF=gi:11345479 /UG=Hs.4746 hypothetical protein FLJ21324 /FL=gb:NM_021941.1 gb:BC003651.1	NM_021941		NP_068760
218023_s_at	5.03E-04	gb:NM_016605.1 /DEF=Homo sapiens putative nuclear protein (LOC51307), mRNA. /FEA=mRNA /GEN=LOC51307 /PROD=putative nuclear protein /DB_XREF=gi:7706138 /UG=Hs.102469 putative nuclear protein /FL=gb:AF251040.1 gb:NM_016605.1	NM_016605		NP_057689

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218030_at	1.26E-04	gb:NM_014030.1 /DEF=Homo sapiens G protein-coupled receptor kinase-interactor 1 (GIT1), mRNA. /FEA=mRNA /GEN=GIT1 /PROD=G protein-coupled receptor kinase-interactor 1 /DB_XREF=gi:7661711 /UG=Hs.318339 G protein-coupled receptor kinase-interactor 1 /FL=gb:AF124490.1 gb:NM_014030.1	NM_014030		NP_054749
218032_at	0.007779	gb:AF070673.1 /DEF=Homo sapiens stannin mRNA, complete cds. /FEA=mRNA /PROD=stannin /DB_XREF=gi:3978241 /UG=Hs.76691 stannin. /FL=gb:AF030196.1 gb:AF070673.1 gb:NM_003498.1 gb:AL161976.1	AF070673		NP_003489
218034_at	0.005852	gb:NM_016068.1 /DEF=Homo sapiens CGI-135 protein (LOC51024), mRNA. /FEA=mRNA /GEN=LOC51024 /PROD=CGI-135 protein /DB_XREF=gi:7705631 /UG=Hs.84344 CGI-135 protein /FL=gb:BC003540.1 gb:AF151893.1 gb:NM_016068.1	NM_016068		NP_057152
218037_at	0.00161	gb:NM_024293.1 /DEF=Homo sapiens hypothetical protein MGC3035 (MGC3035), mRNA. /FEA=mRNA /GEN=MGC3035 /PROD=hypothetical protein MGC3035 /DB_XREF=gi:13236511 /UG=Hs.22412 hypothetical protein MGC3035 /FL=gb:AL136758.1 gb:BC002420.1 gb:NM_024293.1	NM_024293		NP_077269

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
218039_at	0.00161	gb:NM_016359.1 /DEF=Homo sapiens clone HQ0310 PRO0310p1 (LOC51203), mRNA. /FEA=mRNA /GEN=LOC51203 /PROD=clone HQ0310 PRO0310p1 /DB_XREF=gi:7705950 /UG=Hs.279905 clone HQ0310 PRO0310p1 /FL=gb:AF305711.1 gb:BC001308.1 gb:AF290612.1 gb:AF090915.1 gb:NM_016359.1	NM_016359		NP_060924
218041_x_at	1.26E-04	gb:NM_018573.1 /DEF=Homo sapiens hypothetical protein PRO1068 (PRO1068), mRNA. /FEA=mRNA /GEN=PRO1068 /PROD=hypothetical protein PRO1068 /DB_XREF=gi:8924006 /UG=Hs.321158 hypothetical protein PRO1068 /FL=gb:AF116620.1 gb:NM_018573.1	NM_018573		
218043_s_at	0.00228	gb:NM_022461.1 /DEF=Homo sapiens hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2 (FLJ21939), mRNA. /FEA=mRNA /GEN=FLJ21939 /PROD=hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2 /DB_XREF=gi:11968002 /UG=Hs.164478 hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2 /FL=gb:NM_022461.1	NM_022461		NP_071906
218047_at	5.03E-04	gb:NM_024586.1 /DEF=Homo sapiens hypothetical protein FLJ12492 (FLJ12492), mRNA. /FEA=mRNA /GEN=FLJ12492 /PROD=hypothetical protein FLJ12492 /DB_XREF=gi:13375771 /UG=Hs.21938 hypothetical protein FLJ12492 /FL=gb:NM_024586.1	NM_024586		NP_683707

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218049_s_at	0.02141	gb:NM_014078.1 /DEF=Homo sapiens L13 protein (L13), mRNA. /FEA=mRNA /GEN=L13 /PROD=L13 protein /DB_XREF=gi:7662495 /UG=Hs.43946 L13 protein /FL=gb:AF112214.1 gb:NM_014078.1	NM_014078		NP_054797
218057_x_at	0.003171	gb:NM_006067.1 /DEF=Homo sapiens neighbor of COX4 (NOC4), mRNA. /FEA=mRNA /GEN=NOC4 /PROD=neighbor of COX4 /DB_XREF=gi:5174614 /UG=Hs.173162 neighbor of COX4 /FL=gb:BC001472.1 gb:AF005888.1 gb:NM_006067.1	NM_006067		NP_006058
218066_at	0.016934	gb:NM_006598.1 /DEF=Homo sapiens solute carrier family 12 (potassiumchloride transporters), member 7 (SLC12A7), mRNA. /FEA=mRNA /GEN=SLC12A7 /PROD=solute carrier family 12 (potassiumchloridetransporters), member 7 /DB_XREF=gi:5730042 /UG=Hs.172613 solute carrier family 12 (potassiumchloride transporters); member 7 /FL=gb:AF105365.1 gb:NM_006598.1	NM_006598		NP_006589
218068_s_at	0.026891	gb:NM_024836.1 /DEF=Homo sapiens hypothetical protein FLJ22301 (FLJ22301), mRNA. /FEA=mRNA /GEN=FLJ22301 /PROD=hypothetical protein FLJ22301 /DB_XREF=gi:13376246 /UG=Hs.181406 hypothetical protein FLJ22301 /FL=gb:NM_024836.1	NM_024836		NP_079112
218071_s_at	7.58E-04	gb:NM_014160.2 /DEF=Homo sapiens HSPC070 protein (HSPC070), mRNA. /FEA=mRNA /GEN=HSPC070 /PROD=HSPC070 protein /DB_XREF=gi:8850222 /UG=Hs.279474 HSPC070 protein /FL=gb:AF302084.1 gb:AF161555.2 gb:NM_014160.2	NM_014160		NP_054879

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
218077_s_at	0.00476	601063935F1 NIH_MGC_10 Homo sapiens cDNA clone. IMAGE:3450249 5', mRNA sequence.	BE542551		
218078_s_at	1.26E-04	gb:NM_016598.1 /DEF=Homo sapiens DHHC1 protein (LOC51304), mRNA. /FEA=mRNA /GEN=LOC51304 /PROD=DHHC1 protein /DB_XREF=gi:7706132 /UG=Hs.14896 DHHC1 protein /FL=gb:AF247703.1 gb:NM_016598.1	NM_016598		NP_057682
218080_x_at	0.034317	gb:NM_007051.1 /DEF=Homo sapiens Fas (TNFRSF6) associated factor 1 (FAF1), mRNA. /FEA=mRNA /GEN=FAF1 /PROD=Fas (TNFRSF6) associated factor 1 /DB_XREF=gi:5901947 /UG=Hs.25821 Fas (TNFRSF6) associated factor 1 /FL=gb:BC004970.1 gb:AF132938.1 gb:AF106798.1 gb:NM_007051.1	NM_007051		NP_572051
218085_at	0.00161	gb:NM_015961.1 /DEF=Homo sapiens CGI-34 protein (LOC51612), mRNA. /FEA=mRNA /GEN=LOC51612 /PROD=CGI-34 protein /DB_XREF=gi:7706273 /UG=Hs.36237 CGI-34 protein /FL=gb:AF275810.1 gb:AF132968.1 gb:NM_015961.1	NM_015961		
218091_at	1.26E-04	HIV-1 Rev binding protein	AI989512	Hs.171545	NP_004495
218092_s_at	1.26E-04	gb:NM_004504.2 /DEF=Homo sapiens HIV-1 Rev binding protein (HRB), mRNA. /FEA=mRNA /GEN=HRB /PROD=HIV-1 Rev binding protein /DB_XREF=gi:7262381 /UG=Hs.171545 HIV-1 Rev binding protein /FL=gb:NM_004504.2	NM_004504		NP_004495

Gen Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
218093_s_at	7.58E-04	gb:NM_017664.1 /DEF=Homo sapiens hypothetical protein FLJ20093 (FLJ20093), mRNA. /FEA=mRNA /GEN=FLJ20093 /PROD=hypothetical protein FLJ20093 /DB_XREF=gi:8923103 /UG=Hs.172572 hypothetical protein FLJ20093 /FL=gb:NM_017664.1	NM_017664		NP_060134
218095_s_at	0.010205	gb:NM_018475.1 /DEF=Homo sapiens uncharacterized hypothalamus protein HTMP (LOC55858), mRNA. /FEA=mRNA /GEN=LOC55858 /PROD=uncharacterized hypothalamus protein HTMP /DB_XREF=gi:8923860 /UG=Hs.236510 uncharacterized hypothalamus protein HTMP /FL=gb:BC003545.1 gb:AF220188.1 gb:NM_018475.1 gb:AF183409.1	NM_018475		NP_060945
218097_s_at	0.003171	gb:NM_024040.1 /DEF=Homo sapiens hypothetical protein MGC2491 (MGC2491), mRNA. /FEA=mRNA /GEN=MGC2491 /PROD=hypothetical protein MGC2491 /DB_XREF=gi:13128995 /UG=Hs.11270 hypothetical protein MGC2491 /FL=gb:BC000262.1 gb:NM_024040.1	NM_024040		NP_076945
218099_at	0.041973	gb:NM_018469.1 /DEF=Homo sapiens uncharacterized hypothalamus protein HT008 (HT008), mRNA. /FEA=mRNA /GEN=HT008 /PROD=uncharacterized hypothalamus protein HT008 /DB_XREF=gi:8923803 /UG=Hs.16206 uncharacterized hypothalamus protein HT008 /FL=gb:AF220182.1 gb:NM_018469.1	NM_018469		NP_060939

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218101_s_at	3.26E-04	gb:NM_004549.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B14.5b) (NDUFC2), mRNA. /FEA=mRNA /GEN=NDUFC2 /PROD=NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B14.5b) /DB_XREF=gi:4758783 /UG=Hs.193313 NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B14.5b) /FL=gb:AF087659.1 gb:AF070652.1 gb:NM_004549.1	NM_004549		NP_004540
218102_at	0.00161	gb:NM_015954.1 /DEF=Homo sapiens CGI-26 protein (LOC51071), mRNA. /FEA=mRNA /GEN=LOC51071 /PROD=CGI-26 protein /DB_XREF=gi:7705717 /UG=Hs.24332 CGI-26 protein /FL=gb:AF132960.1 gb:NM_015954.1	NM_015954		NP_057038
218107_at	2.05E-04	gb:NM_025160.1 /DEF=Homo sapiens hypothetical protein FLJ21016 (FLJ21016), mRNA. /FEA=mRNA /GEN=FLJ21016 /PROD=hypothetical protein FLJ21016 /DB_XREF=gi:13376760 /UG=Hs.289069 hypothetical protein FLJ21016 /FL=gb:NM_025160.1	NM_025160		NP_079436
218108_at	2.05E-04	gb:NM_018108.1 /DEF=Homo sapiens hypothetical protein FLJ10483 (FLJ10483), mRNA. /FEA=mRNA /GEN=FLJ10483 /PROD=hypothetical protein FLJ10483 /DB_XREF=gi:8922451 /UG=Hs.6877 hypothetical protein FLJ10483 /FL=gb:NM_018108.1	NM_018108		NP_786924

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218109_s_at	1.26E-04	gb:NM_022736.1 /DEF=Homo sapiens hypothetical protein FLJ14153 (FLJ14153), mRNA. /FEA=mRNA /GEN=FLJ14153 /PROD=hypothetical protein FLJ14153 /DB_XREF=gi:12232392 /UG=Hs.7503 hypothetical protein FLJ14153 /FL=gb:NM_022736.1 gb:AB014732.1	NM_022736		NP_073573
218111_s_at	0.002313	gb:NM_018686.1 /DEF=Homo sapiens CMP-N-acetylneuraminic acid synthase (LOC55907), mRNA. /FEA=mRNA /GEN=LOC55907 /PROD=CMP-N-acetylneuraminic acid synthase /DB_XREF=gi:8923899 /UG=Hs.12492 CMP-N-acetylneuraminic acid synthase /FL=gb:AF271388.1 gb:NM_018686.1	NM_018686		NP_061156
218113_at	5.03E-04	gb:NM_013390.1 /DEF=Homo sapiens transmembrane protein 2 (TMEM2), mRNA. /FEA=mRNA /GEN=TMEM2 /PROD=transmembrane protein 2 /DB_XREF=gi:7019554 /UG=Hs.160417 transmembrane protein 2 /FL=gb:AF137030.1 gb:NM_013390.1	NM_013390		NP_037522
218116_at	0.026891	gb:NM_016520.1 /DEF=Homo sapiens hepatocellular carcinoma-associated antigen 59 (LOC51759), mRNA. /FEA=mRNA /GEN=LOC51759 /PROD=hepatocellular carcinoma-associated antigen 59. /DB_XREF=gi:7706556 /UG=Hs.278429 hepatocellular carcinoma-associated antigen 59 /FL=gb:AF151054.1 gb:AF218421.1 gb:NM_016482.1 gb:NM_016520.1	NM_016520		NP_057604

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Prot in Accession No.
218117_at	0.00228	gb:NM_014248.1 /DEF=Homo sapiens ring-box 1 (RBX1), mRNA. /FEA=mRNA /GEN=RBX1 /PROD=ring-box 1 /DB_XREF=gi:7657507 /UG=Hs.279919 ring-box 1 /FL=gb:BC001466.1 gb:AF140598.1 gb:AF142059.1 gb:NM_014248.1	NM_014248		NP_055063
218118_s_at	0.010061	gb:NM_006327.1 /DEF=Homo sapiens translocase of inner mitochondrial membrane 23 (yeast) homolog (TIM23), mRNA. /FEA=mRNA /GEN=TIM23 /PROD=translocase of inner mitochondrial membrane 23(yeast) homolog /DB_XREF=gi:5454121 /UG=Hs.283684 translocase of inner mitochondrial membrane 23 (yeast) homolog /FL=gb:AF030162.1 gb:NM_006327.1	NM_006327		NP_006318
218119_at	0.033381	gb:NM_006327.1 /DEF=Homo sapiens translocase of inner mitochondrial membrane 23 (yeast) homolog (TIM23), mRNA. /FEA=mRNA /GEN=TIM23 /PROD=translocase of inner mitochondrial membrane 23(yeast) homolog /DB_XREF=gi:5454121 /UG=Hs.283684 translocase of inner mitochondrial membrane 23 (yeast) homolog /FL=gb:AF030162.1 gb:NM_006327.1	NM_006327		NP_006318
218122_s_at	0.013088	gb:NM_021627.1 /DEF=Homo sapiens sentrin-specific protease (SENP2), mRNA. /FEA=mRNA /GEN=SENP2 /PROD=sentrin-specific protease /DB_XREF=gi:11055993 /UG=Hs.3355 sentrin-specific protease /FL=gb:AF151697.2 gb:NM_021627.1	NM_021627		NP_067640

Gene Identifier	p-value	D scription	Gene Accession No.	Unigen Accession No.	Protein Accession No.
218124_at	7.58E-04	gb:NM_017750.1 /DEF=Homo sapiens hypothetical protein FLJ20296 (FLJ20296), mRNA /FEA=mRNA /GEN=FLJ20296 /PROD=hypothetical protein FLJ20296 /DB_XREF=gi:8923274 /UG=Hs.6603 hypothetical protein FLJ20296 /FL=gb:NM_017750.1	NM_017750		NP_060220
218129_s_at	0.001078	gb:NM_006166.2 /DEF=Homo sapiens nuclear transcription factor Y, beta (NFYB), mRNA /FEA=mRNA /GEN=NFYB /PROD=nuclear transcription factor Y, beta /DB_XREF=gi:11496976 /UG=Hs.84928 nuclear transcription factor Y, beta /FL=gb:NM_006166.2 gb:BC005316.1 gb:BC005317.1 gb:L06145.1	NM_006166		NP_006157
218130_at	1.26E-04	gb:NM_024510.1 /DEF=Homo sapiens hypothetical protein MGC4368 (MGC4368), mRNA /FEA=mRNA /GEN=MGC4368 /PROD=hypothetical protein MGC4368 /DB_XREF=gi:13375647 /UG=Hs.9732 hypothetical protein MGC4368 /FL=gb:BC003595.1 gb:BC004171.1 gb:NM_024510.1	NM_024510		NP_078786
218131_s_at	0.004341	gb:NM_017660.1 /DEF=Homo sapiens hypothetical protein FLJ20085 (FLJ20085), mRNA /FEA=mRNA /GEN=FLJ20085 /PROD=hypothetical protein FLJ20085 /DB_XREF=gi:8923093 /UG=Hs.118964 hypothetical protein FLJ20085 /FL=gb:NM_017660.1	NM_017660		NP_060130

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218134_s_at	0.010205	gb:NM_018047.1 /DEF=Homo sapiens hypothetical protein FLJ10290 (FLJ10290), mRNA. /FEA=mRNA /GEN=FLJ10290 /PROD=hypothetical protein FLJ10290 /DB_XREF=gi:8922327 /UG=Hs.25516 hypothetical protein FLJ10290 /FL=gb:AL136933.1 gb:BC003402.1 gb:NM_018047.1	NM_018047		NP_060517
218135_at	0.00228	gb:NM_016570.1 /DEF=Homo sapiens CDA14 (LOC51290), mRNA. /FEA=mRNA /GEN=LOC51290 /PROD=CDA14 /DB_XREF=gi:7706104 /UG=Hs.26813 CDA14 /FL=gb:BC000887.1 gb:AF216751.1 gb:NM_016570.1 gb:AF183410.1	NM_016570		NP_057654
218137_s_at	5.03E-04	gb:NM_021940.1 /DEF=Homo sapiens hypothetical protein FLJ13159 (FLJ13159), mRNA. /FEA=mRNA /GEN=FLJ13159 /PROD=hypothetical protein FLJ13159 /DB_XREF=gi:11345477 /UG=Hs.4273 hypothetical protein FLJ13159 /FL=gb:NM_021940.1	NM_021940		NP_068759
218138_at	0.005852	gb:NM_018848.1 /DEF=Homo sapiens McKusick-Kaufman syndrome (MKKS), mRNA. /FEA=mRNA /GEN=MKKS /PROD=McKusick-Kaufman syndrome protein /DB_XREF=gi:9055271 /UG=Hs.46743 McKusick-Kaufman syndrome /FL=gb:AF275813.1 gb:AF221992.1 gb:AF221993.1 gb:NM_018848.1	NM_018848		NP_740754

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218139_s_at	0.010205	gb:NM_018229.1 /DEF=Homo sapiens hypothetical protein FLJ10813 (FLJ10813), mRNA. /FEA=mRNA /GEN=FLJ10813 /PROD=hypothetical protein FLJ10813 /DB_XREF=gi:8922687 /UG=Hs.106210 hypothetical protein FLJ10813 /FL=gb:AL136685.1 gb:NM_018229.1	NM_018229		NP_060699
218142_s_at	0.00161	gb:NM_016302.1 /DEF=Homo sapiens protein x 0001 (LOC51185), mRNA. /FEA=mRNA /GEN=LOC51185 /PROD=protein x 0001 /DB_XREF=gi:10047097 /UG=Hs.18925 protein x 0001 /FL=gb:NM_016302.1 gb:AF117230.1	NM_016302		NP_057386
218157_x_at	0.00161	gb:NM_020239.2 /DEF=Homo sapiens small protein effector 1 of Cdc42 (SPEC1), mRNA. /FEA=mRNA /GEN=SPEC1 /PROD=small protein effector 1 of Cdc42 /DB_XREF=gi:12965169 /UG=Hs.22065 small protein effector 1 of Cdc42 /FL=gb:AF187845.2 gb:NM_020239.2	NM_020239		NP_064624
218158_s_at	0.036254	gb:NM_012096.1 /DEF=Homo sapiens adaptor protein containing pH domain, PTB domain and leucine zipper motif (APPL), mRNA. /FEA=mRNA /GEN=APPL /PROD=adaptor protein containing pH domain, PTB domain and leucine zipper motif /DB_XREF=gi:6912241 /UG=Hs.27413 adaptor protein containing pH domain, PTB domain and leucine zipper motif /FL=gb:AF169797.1 gb:NM_012096.1	NM_012096		NP_036228

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
218159_at	0.007258	gb:NM_023935.1 /DEF=Homo sapiens hypothetical protein MGC2592 (MGC2592), mRNA. /FEA=mRNA /GEN=MGC2592 /PROD=hypothetical protein MGC2592 /DB_XREF=gi:13027601 /UG=Hs.28393 hypothetical protein MGC2592 /FL=gb:BC000643.1 gb:NM_023935.1	NM_023935		NP_076424
218163_at	2.05E-04	gb:NM_014060.1 /DEF=Homo sapiens MCT-1 protein (MCT-1), mRNA. /FEA=mRNA /GEN=MCT-1 /PROD=MCT-1 protein /DB_XREF=gi:7662501 /UG=Hs.102696 MCT-1 protein /FL=gb:BC001013.1 gb:AB034206.1 gb:NM_014060.1	NM_014060		NP_054779
218167_at	0.033381	gb:NM_016627.1 /DEF=Homo sapiens hypothetical protein (LOC51321), mRNA. /FEA=mRNA /GEN=LOC51321 /PROD=hypothetical protein /DB_XREF=gi:7706167 /UG=Hs.268122 hypothetical protein /FL=gb:AF208856.1 gb:NM_016627.1	NM_016627		NP_057711
218168_s_at	0.007779	gb:NM_020247.1 /DEF=Homo sapiens hypothetical protein, clone Telethon(Italy_B41)_Strait02270_F L142 (LOC56997), mRNA. /FEA=mRNA /GEN=LOC56997 /PROD=hypothetical protein, clone Telethon(Italy_B41)_Strait02270_FL142 /DB_XREF=gi:9910387 /UG=Hs.273186 hypothetical protein, clone Telethon(Italy_B41)_Strait02270_F L142 /FL=gb:BC005171.1 gb:NM_020247.1	NM_020247		NP_064632

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
218171_at	2.05E-04	gb:AF195514.1 /DEF=Homo sapiens VPS4-2 ATPase (VPS42) mRNA, complete cds. /FEA=mRNA /GEN=VPS42 /PROD=VPS4-2 ATPase /DB_XREF=gi:11225484 /UG=Hs.126550 suppressor of K+ transport defect 1 /FL=gb:AF195514.1 gb:AF038960.1 gb:NM_004869.1	AF195514		NP_004860
218177_at	0.016934	CHMP1.5 protein	AA293502	Hs.380932	NP_065145
218178_s_at	0.02145	gb:NM_020412.1 /DEF=Homo sapiens CHMP1.5 protein (CHMP1.5), mRNA. /FEA=mRNA /GEN=CHMP1.5 /PROD=CHMP1.5 protein /DB_XREF=gi:9966900 /UG=Hs.42733 CHMP1.5 protein /FL=gb:AF281064.1 gb:NM_020412.1	NM_020412		NP_065145
218179_s_at	0.02145	gb:NM_021942.1 /DEF=Homo sapiens hypothetical protein FLJ12716 (FLJ12716), mRNA. /FEA=mRNA /GEN=FLJ12716 /PROD=hypothetical protein FLJ12716 /DB_XREF=gi:11345481 /UG=Hs.5354 hypothetical protein FLJ12716 /FL=gb:NM_021942.1 gb:AL136752.1	NM_021942		NP_068761
218181_s_at	1.26E-04	gb:NM_017792.1 /DEF=Homo sapiens hypothetical protein FLJ20373 (FLJ20373), mRNA. /FEA=mRNA /GEN=FLJ20373 /PROD=hypothetical protein FLJ20373 /DB_XREF=gi:8923351 /UG=Hs.6631 hypothetical protein FLJ20373 /FL=gb:NM_017792.1	NM_017792		NP_060262
218185_s_at	0.013223	gb:NM_018120.1 /DEF=Homo sapiens hypothetical protein FLJ10511 (FLJ10511), mRNA. /FEA=mRNA /GEN=FLJ10511 /PROD=hypothetical protein FLJ10511 /DB_XREF=gi:8922478 /UG=Hs.106768 hypothetical protein FLJ10511 /FL=gb:NM_018120.1	NM_018120		NP_060590

Gene Identifier	p-value	Description	Gene Accession No.	UniProt Accession No.	Protein Accession No.
218187_s_at	0.02145	gb:NM_023080.1 /DEF=Homo sapiens hypothetical protein FLJ20989 (FLJ20989), mRNA. /FEA=mRNA /GEN=FLJ20989 /PROD=hypothetical protein FLJ20989 /DB_XREF=gi:12751496 /UG=Hs.169615 hypothetical protein FLJ20989 /FL=gb:NM_023080.1	NM_023080		NP_075568
218189_s_at	3.26E-04	gb:NM_018946.2 /DEF=Homo sapiens N-acetylneuraminic acid phosphate synthase, sialic acid synthase (SAS), mRNA. /FEA=mRNA /GEN=SAS /PROD=N-acetylneuraminic acid phosphate synthase /DB_XREF=gi:12056472 /UG=Hs.274424 N-acetylneuraminic acid phosphate synthase, sialic acid synthase /FL=gb:NM_018946.2 gb:BC000008.1 gb:AF257466.1	NM_018946		NP_061819
218191_s_at	0.003171	gb:NM_018368.1 /DEF=Homo sapiens hypothetical protein FLJ11240 (FLJ11240), mRNA. /FEA=mRNA /GEN=FLJ11240 /PROD=hypothetical protein FLJ11240 /DB_XREF=gi:8922955 /UG=Hs.301308 hypothetical protein FLJ11240 /FL=gb:AF113224.1 gb:AF211480.1 gb:NM_018368.1	NM_018368		NP_060838
218195_at	1.26E-04	gb:NM_024573.1 /DEF=Homo sapiens hypothetical protein FLJ12910 (FLJ12910), mRNA. /FEA=mRNA /GEN=FLJ12910 /PROD=hypothetical protein FLJ12910 /DB_XREF=gi:13375745 /UG=Hs.15929 hypothetical protein FLJ12910 /FL=gb:NM_024573.1	NM_024573		NP_078849

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Acc ssion No.
218196_at	3.26E-04	gb:NM_014028.1 /DEF=Homo sapiens HSPC019 protein (HSPC019), mRNA. /FEA=mRNA /GEN=HSPC019 /PROD=HSPC019 protein /DB_XREF=gi:7661737 /UG=Hs.163724 HSPC019 protein /FL=gb:AF077205.1 gb:NM_014028.1	NM_014028		NP_054747
218202_x_at	0.00161	gb:NM_022915.1 /DEF=Homo sapiens hypothetical protein FLJ12701 (FLJ12701), mRNA. /FEA=mRNA /GEN=FLJ12701 /PROD=hypothetical protein FLJ12701 /DB_XREF=gi:12597660 /UG=Hs.203559 hypothetical protein FLJ12701 /FL=gb:NM_022915.1	NM_022915		NP_075066
218208_at	3.26E-04	gb:NM_025078.1 /DEF=Homo sapiens hypothetical protein FLJ22378 (FLJ22378), mRNA. /FEA=mRNA /GEN=FLJ22378 /PROD=hypothetical protein FLJ22378 /DB_XREF=gi:13376629 /UG=Hs.288284 hypothetical protein FLJ22378 /FL=gb:NM_025078.1	NM_025078		NP_079354
218209_s_at	3.26E-04	gb:NM_018170.1 /DEF=Homo sapiens hypothetical protein FLJ10656 (FLJ10656), mRNA. /FEA=mRNA /GEN=FLJ10656 /PROD=hypothetical protein FLJ10656 /DB_XREF=gi:8922574 /UG=Hs.300906 hypothetical protein FLJ10656 /FL=gb:NM_018170.1	NM_018170		NP_060640

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218213_s_at	0.00161	gb:NM_014206.1 /DEF=Homo sapiens chromosome 11 open reading frame 10 (C11orf10), mRNA. /FEA=mRNA /GEN=C11orf10 /PROD=chromosome 11 open reading frame 10 /DB_XREF=gi:7656933 /UG=Hs.90918 chromosome 11 open reading frame 10 /FL=gb:BC002750.1 gb:AF070661.1 gb:AF086763.1 gb:NM_014206.1	NM_014206		NP_055021
218217_at	1.26E-04	gb:NM_021626.1 /DEF=Homo sapiens serine carboxypeptidase 1 precursor protein (HSCP1), mRNA. /FEA=mRNA /GEN=HSCP1 /PROD=serine carboxypeptidase 1 precursor protein /DB_XREF=gi:11055991 /UG=Hs.106747 serine carboxypeptidase 1 precursor protein /FL=gb:AF282618.1 gb:NM_021626.1 gb:AF113214.1 gb:AF265441.1	NM_021626		NP_067639
218218_at	1.26E-04	gb:NM_018171.1 /DEF=Homo sapiens hypothetical protein FLJ10659 (FLJ10659), mRNA. /FEA=mRNA /GEN=FLJ10659 /PROD=hypothetical protein FLJ10659 /DB_XREF=gi:8922576 /UG=Hs.107882 hypothetical protein FLJ10659 /FL=gb:NM_018171.1	NM_018171		NP_060641
218221_at	1.26E-04	ESTs	AL042842	Hs.166172	NP_848514
218223_s_at	0.001116	gb:NM_016274.1 /DEF=Homo sapiens CK2 interacting protein 1; HQ0024c protein (LOC51177), mRNA. /FEA=mRNA /GEN=LOC51177 /PROD=CK2 interacting protein 1; HQ0024c protein /DB_XREF=gi:7705918 /UG=Hs.173380 CK2 interacting protein 1; HQ0024c protein /FL=gb:AF291105.1 gb:AF073836.1 gb:NM_016274.1 gb:AF168676.1	NM_016274		NP_057358

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218226_s_at	0.010205	gb:NM_004547.2 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD, B15) (NDUFB4), mRNA. /FEA=mRNA /GEN=NDUFB4 /PROD=NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD, B15) /DB_XREF=gi:6041668 /UG=Hs.227750. NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD, B15) /FL=gb:BC000855.1 gb:AF044957.1 gb:NM_004547.2	NM_004547		NP_004538
218228_s_at	0.005852	gb:NM_025235.1 /DEF=Homo sapiens tankyrase 2 (TNKL), mRNA. /FEA=mRNA /GEN=TNKL /PROD=tankyrase 2 /DB_XREF=gi:13376841 /UG=Hs.280776 tankyrase 2 /FL=gb:AF264912.1 gb:AF329696.1 gb:NM_025235.1 gb:AF342982.1	NM_025235		NP_079511
218237_s_at	0.001116	gb:NM_030674.1 /DEF=Homo sapiens amino acid transporter system A1 (ATA1), mRNA. /FEA=mRNA /GEN=ATA1 /PROD=amino acid transporter system A1 /DB_XREF=gi:13492978 /UG=Hs.18272 amino acid transporter system A1 /FL=gb:AF271070.1 gb:NM_030674.1	NM_030674		NP_109599
218243_at	5.03E-04	gb:NM_025158.1 /DEF=Homo sapiens hypothetical protein FLJ22251 (FLJ22251), mRNA. /FEA=mRNA /GEN=FLJ22251 /PROD=hypothetical protein FLJ22251 /DB_XREF=gi:13376756 /UG=Hs.289064 hypothetical protein FLJ22251 /FL=gb:NM_025158.1	NM_025158		NP_079434

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218249_at	0.033381	gb:NM_022494.1 /DEF=Homo sapiens hypothetical protein FLJ21952 (FLJ21952), mRNA. /FEA=mRNA /GEN=FLJ21952 /PROD=hypothetical protein FLJ21952 /DB_XREF=gi:11968052 /UG=Hs.22353 hypothetical protein FLJ21952 /FL=gb:NM_022494.1	NM_022494		NP_071939
218250_s_at	0.004341	gb:NM_013354.2 /DEF=Homo sapiens CCR4-NOT transcription complex, subunit 7 (CNOT7), mRNA. /FEA=mRNA /GEN=CNOT7 /PROD=CCR4-NOT transcription complex, subunit 7 /DB_XREF=gi:10518495 /UG=Hs.226318 CCR4-NOT transcription complex, subunit 7 /FL=gb:NM_013354.2 gb:L46722.1	NM_013354		NP_473367
218251_at	2.05E-04	gb:NM_021242.1 /DEF=Homo sapiens hypothetical protein STRAIT11499 (STRAIT11499), mRNA. /FEA=mRNA /GEN=STRAIT11499 /PROD=hypothetical protein STRAIT11499 /DB_XREF=gi:10864050 /UG=Hs.236556 hypothetical protein STRAIT11499 /FL=gb:NM_021242.1	NM_021242		NP_067065
218252_at	3.26E-04	gb:NM_018204.1 /DEF=Homo sapiens cytoskeleton associated protein 2 (CKAP2), mRNA. /FEA=mRNA /GEN=CKAP2 /PROD=cytoskeleton associated protein 2 /DB_XREF=gi:8922641 /UG=Hs.24641 cytoskeleton associated protein 2 /FL=gb:AL136848.1 gb:NM_018204.1	NM_018204		NP_060674

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218253_s_at	0.028893	gb:NM_006893.1 /DEF=Homo sapiens ligatin (LGTN), mRNA. /FEA=mRNA /GEN=LGTN /PROD=ligatin /DB_XREF=gi:11386190 /UG=Hs.274151 ligatin /FL=gb:NM_006893.1 gb:AF220417.2	NM_006893		NP_008824
218254_s_at	2.05E-04	gb:NM_016103.1 /DEF=Homo sapiens GTP-binding protein Sara (LOC51128), mRNA. /FEA=mRNA /GEN=LOC51128 /PROD=GTP-binding protein Sara /DB_XREF=gi:7705826 /UG=Hs.279582 GTP-binding protein Sara /FL=gb:AF092130.1 gb:NM_016103.1	NM_016103		NP_057187
218255_s_at	0.041055	gb:NM_022452.1 /DEF=Homo sapiens hypothetical protein FLJ11618 (FLJ11618), mRNA. /FEA=mRNA /GEN=FLJ11618 /PROD=hypothetical protein FLJ11618 /DB_XREF=gi:11967986 /UG=Hs.77735 hypothetical protein FLJ11618 /FL=gb:NM_022452.1	NM_022452		NP_071897
218262_at	7.58E-04	gb:NM_022762.1 /DEF=Homo sapiens hypothetical protein FLJ22318 (FLJ22318), mRNA. /FEA=mRNA /GEN=FLJ22318 /PROD=hypothetical protein FLJ22318 /DB_XREF=gi:12232432 /UG=Hs.22753 hypothetical protein FLJ22318 /FL=gb:NM_022762.1	NM_022762		NP_073599
218268_at	0.013223	gb:NM_022771.1 /DEF=Homo sapiens hypothetical protein FLJ12085 (FLJ12085), mRNA. /FEA=mRNA /GEN=FLJ12085 /PROD=hypothetical protein FLJ12085 /DB_XREF=gi:12232450 /UG=Hs.48827 hypothetical protein FLJ12085 /FL=gb:NM_022771.1	NM_022771		NP_073608

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Prot in Accession No.
218271_s_at	3.26E-04	gb:NM_018622.1 /DEF=Homo sapiens presenilins associated rhomboid-like protein (PRO2207), mRNA. /FEA=mRNA /GEN=PRO2207 /PROD=presenilins associated rhomboid-like protein /DB_XREF=gi:8924133 /UG=Hs.13094 presenilins associated rhomboid-like protein /FL=gb:AF197937.1 gb:AF116692.1 gb:NM_018622.1	NM_018622		NP_061092
218273_s_at	7.58E-04	gb:NM_018444.1 /DEF=Homo sapiens pyruvate dehydrogenase phosphatase (PDP), mRNA. /FEA=mRNA /GEN=PDP /PROD=pyruvate dehydrogenase phosphatase /DB_XREF=gi:8923959 /UG=Hs.22265 pyruvate dehydrogenase phosphatase /FL=gb:AF155661.1 gb:NM_018444.1	NM_018444		NP_060914
218277_s_at	0.010205	gb:NM_024612.1 /DEF=Homo sapiens hypothetical protein FLJ22060 (FLJ22060), mRNA. /FEA=mRNA /GEN=FLJ22060 /PROD=hypothetical protein FLJ22060 /DB_XREF=gi:13375824 /UG=Hs.29403 hypothetical protein FLJ22060 /FL=gb:NM_024612.1	NM_024612		NP_078888
218280_x_at	5.03E-04	gb:NM_003516.1 /DEF=Homo sapiens H2A histone family, member O (H2AFO), mRNA. /FEA=mRNA /GEN=H2AFO /PROD=H2A histone family, member O /DB_XREF=gi:4504250 /UG=Hs.795 H2A histone family, member O /FL=gb:BC001629.1 gb:L19779.1 gb:NM_003516.1	NM_003516		NP_003507

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
218286_s_at	0.003171	gb:NM_014245.1 /DEF=Homo sapiens ring finger protein 7 (RNF7), mRNA. /FEA=mRNA /GEN=RNF7 /PROD=ring finger protein 7 /DB_XREF=gi:7657521 /UG=Hs.14084 ring finger protein 7 /FL=gb:AF092878.1 gb:AF164679.1 gb:NM_014245.1	NM_014245		NP_055060
218291_at	7.58E-04	gb:NM_014017.1 /DEF=Homo sapiens HSPC003 protein (HSPC003), mRNA. /FEA=mRNA /GEN=HSPC003 /PROD=HSPC003 protein /DB_XREF=gi:7661727 /UG=Hs.25635 HSPC003 protein /FL=gb:AF070659.1 gb:NM_014017.1	NM_014017		NP_054736
218301_at	0.010205	gb:NM_018226.1 /DEF=Homo sapiens arginyl aminopeptidase (aminopeptidase B)-like 1 (RNPEPL1), mRNA. /FEA=mRNA /GEN=RNPEPL1 /PROD=arginyl aminopeptidase (aminopeptidase B)-like1 /DB_XREF=gi:8922680 /UG=Hs.5345 arginyl aminopeptidase (aminopeptidase B)-like 1 /FL=gb:AF300795.1 gb:NM_018226.1	NM_018226		NP_060696
218303_x_at	1.26E-04	gb:NM_016618.1 /DEF=Homo sapiens hypothetical protein (LOC51315), mRNA. /FEA=mRNA /GEN=LOC51315 /PROD=hypothetical protein /DB_XREF=gi:7706155 /UG=Hs.5721 hypothetical protein /FL=gb:AF208845.1 gb:AF217520.1 gb:NM_016618.1	NM_016618		NP_057702
218304_s_at	1.26E-04	gb:NM_022776.1 /DEF=Homo sapiens hypothetical protein FLJ13164 (FLJ13164), mRNA. /FEA=mRNA /GEN=FLJ13164 /PROD=hypothetical protein FLJ13164 /DB_XREF=gi:12232460 /UG=Hs.61260 hypothetical protein FLJ13164 /FL=gb:NM_022776.1	NM_022776		NP_073613

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218306_s_at	0.013223	gb:NM_003922.1 /DEF=Homo sapiens hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1 (HERC1), mRNA. /FEA=mRNA /GEN=HERC1 /PROD=guanine nucleotide exchange factor p532 /DB_XREF=gi:4557025 /UG=Hs.76127 hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1 /FL=gb:U50078.1 gb:NM_003922.1	NM_003922		NP_003913
218308_at	0.02145	gb:NM_006342.1 /DEF=Homo sapiens transforming, acidic coiled-coil containing protein 3 (TACC3), mRNA. /FEA=mRNA /GEN=TACC3 /PROD=transforming acidic coiled-coil containing protein 3 /DB_XREF=gi:5454101 /UG=Hs.104019 transforming, acidic coiled-coil containing protein 3 /FL=gb:AF093543.1 gb:NM_006342.1	NM_006342		NP_006333
218310_at	2.05E-04	gb:NM_014504.1 /DEF=Homo sapiens putative Rab5 GDPGTP exchange factor homologue (RABEX5), mRNA. /FEA=mRNA /GEN=RABEX5 /PROD=putative Rab5 GDPGTP exchange factor homologue /DB_XREF=gi:7657495 /UG=Hs.187660 putative Rab5 GDPGTP exchange factor homologue /FL=gb:NM_014504.1	NM_014504		NP_055319

Gene Identifier	p-value	D scription	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218313_s_at	2.05E-04	gb:NM_017423.1 /DEF=Homo sapiens UDP-N-acetyl-alpha-D-galactosamine polypeptide N-acetylgalactosaminyltransferase 7 (GalNAc-T7) (GALNT7); mRNA. /FEA=mRNA /GEN=GALNT7 /PROD=polypeptide N-acetylgalactosaminyltransferase 7 /DB_XREF=gi:8393408 /UG=Hs.246315 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (GalNAc-T7) /FL=gb:NM_017423.1	NM_017423		NP_059119
218318_s_at	0.04734	gb:NM_016231.1 /DEF=Homo sapiens nemo-like kinase (LOC51701), mRNA. /FEA=mRNA /GEN=LOC51701 /PROD=nemo-like kinase /DB_XREF=gi:7706444 /UG=Hs.3532 nemo-like kinase /FL=gb:AF180819.1 gb:AF197898.1 gb:NM_016231.1	NM_016231		NP_057315
218323_at	0.001116	gb:NM_018307.1 /DEF=Homo sapiens hypothetical protein FLJ11040 (FLJ11040), mRNA. /FEA=mRNA /GEN=FLJ11040 /PROD=hypothetical protein FLJ11040 /DB_XREF=gi:8922837 /UG=Hs.14202 hypothetical protein FLJ11040 /FL=gb:NM_018307.1	NM_018307		NP_060777
218332_at	8.94E-04	gb:NM_018476.1 /DEF=Homo sapiens uncharacterized hypothalamus protein HBEX2 (HBEX2), mRNA. /FEA=mRNA /GEN=HBEX2 /PROD=uncharacterized hypothalamus protein HBEX2 /DB_XREF=gi:8923715 /UG=Hs.283719 uncharacterized hypothalamus protein HBEX2 /FL=gb:AF220189.1 gb:NM_018476.1 gb:AF183416.1 gb:AF237783.1	NM_018476		NP_060946

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218333_at	0.007779	gb:NM_016041.1 /DEF=Homo sapiens CGI-101 protein (LOC51009), mRNA. /FEA=mRNA /GEN=LOC51009 /PROD=CGI-101 protein /DB_XREF=gi:7705603 /UG=Hs.286131 CGI-101 protein /FL=gb:AF132289.1 gb:AF151859.1 gb:NM_016041.1 gb:AF242523.1	NM_016041		NP_057125
218334_at	0.00228	gb:NM_025075.1 /DEF=Homo sapiens hypothetical protein FLJ23445 (FLJ23445), mRNA. /FEA=mRNA /GEN=FLJ23445 /PROD=hypothetical protein FLJ23445 /DB_XREF=gi:13376622 /UG=Hs.288151 hypothetical protein FLJ23445 /FL=gb:NM_025075.1	NM_025075		NP_079351
218341_at	0.004341	gb:NM_024664.1 /DEF=Homo sapiens hypothetical protein FLJ11838 (FLJ11838), mRNA. /FEA=mRNA /GEN=FLJ11838 /PROD=hypothetical protein FLJ11838 /DB_XREF=gi:13375918 /UG=Hs.72531 hypothetical protein FLJ11838 /FL=gb:NM_024664.1	NM_024664		NP_078940
218343_s_at	0.026891	gb:NM_012086.1 /DEF=Homo sapiens general transcription factor IIIC, polypeptide 3 (102kD) (GTF3C3), mRNA. /FEA=mRNA /GEN=GTF3C3 /PROD=general transcription factor IIIC, polypeptide 3(102kD) /DB_XREF=gi:6912397 /UG=Hs.90847 general transcription factor IIIC, polypeptide 3 (102kD) /FL=gb:AF133123.1 gb:NM_012086.1	NM_012086		NP_036218

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218344_s_at	0.013223	gb:NM_018254.1 /DEF=Homo sapiens hypothetical protein FLJ10876 (FLJ10876), mRNA. /FEA=mRNA /GEN=FLJ10876 /PROD=hypothetical protein FLJ10876 /DB_XREF=gi:8922732 /UG=Hs.94042 hypothetical protein FLJ10876 /FL=gb:NM_018254.1	NM_018254		NP_060724
218345_at	0.02049	gb:NM_018487.1 /DEF=Homo sapiens hepatocellular carcinoma-associated antigen 112 (HCA112), mRNA. /FEA=mRNA /GEN=HCA112 /PROD=hepatocellular carcinoma-associated antigen 112 /DB_XREF=gi:8923717 /UG=Hs.12126 hepatocellular carcinoma-associated antigen 112 /FL=gb:AF258340.1 gb:NM_018487.1	NM_018487		NP_060957
218348_s_at	2.05E-04	gb:NM_014153.1 /DEF=Homo sapiens HSPC055 protein (HSPC055), mRNA. /FEA=mRNA /GEN=HSPC055 /PROD=HSPC055 protein /DB_XREF=gi:7661761 /UG=Hs.179898 HSPC055 protein /FL=gb:AF161540.1 gb:NM_014153.1	NM_014153		NP_054872
218350_s_at	0.002743	gb:NM_015895.1 /DEF=Homo sapiens geminin (LOC51053), mRNA. /FEA=mRNA /GEN=LOC51053 /PROD=geminin /DB_XREF=gi:7705681 /UG=Hs.234896 geminin /FL=gb:AF067855.1 gb:NM_015895.1	NM_015895		NP_056979
218351_at	0.026891	gb:NM_017845.1 /DEF=Homo sapiens hypothetical protein FLJ20502 (FLJ20502), mRNA. /FEA=mRNA /GEN=FLJ20502 /PROD=hypothetical protein FLJ20502 /DB_XREF=gi:8923457 /UG=Hs.23956 hypothetical protein FLJ20502 /FL=gb:AF182421.1 gb:NM_017845.1	NM_017845		NP_060315

Gene Identifier	p-value	Description	Gene Accession No.	Unig n Accession No.	Protein Accession No.
218352_at	3.26E-04	gb:NM_018191.1 /DEF=Homo sapiens hypothetical protein FLJ10716 (FLJ10716); mRNA. /FEA=mRNA /GEN=FLJ10716 /PROD=hypothetical protein FLJ10716 /DB_XREF=gi:8922616 /UG=Hs.24129 hypothetical protein FLJ10716 /FL=gb:NM_018191.1	NM_018191		NP_060661
218354_at	1.26E-04	gb:NM_016209.1 /DEF=Homo sapiens unknown (LOC51693); mRNA. /FEA=mRNA /GEN=LOC51693 /PROD=unknown /DB_XREF=gi:7706428 /UG=Hs.27445 unknown /FL=gb:AF089106.1 gb:AF161524.1 gb:NM_016209.1	NM_016209		NP_057293
218357_s_at	2.05E-04	gb:NM_012459.1 /DEF=Homo sapiens translocase of inner mitochondrial membrane 8 (yeast) homolog B (TIMM8B); mRNA. /FEA=mRNA /GEN=TIMM8B /PROD=translocase of inner mitochondrial membrane 8(yeast) homolog B /DB_XREF=gi:6912711 /UG=Hs.279915 translocase of inner mitochondrial membrane 8 (yeast) homolog B /FL=gb:AF150087.1 gb:AF152350.1 gb:NM_012459.1	NM_012459		NP_036591
218360_at	1.26E-04	gb:NM_020673.1 /DEF=Homo sapiens RAB22A, member RAS oncogene family (RAB22A); mRNA. /FEA=mRNA /GEN=RAB22A /PROD=RAB22A, member RAS oncogene family /DB_XREF=gi:10190713 /UG=Hs.288968 RAB22A, member RAS oncogene family /FL=gb:NM_020673.1 gb:AF091034.2	NM_020673		NP_065724

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
218364_at	7.58E-04	gb:NM_017724.1 /DEF=Homo sapiens leucine rich repeat (in FLII) interacting protein 2 (LRRFIP2), mRNA. /FEA=mRNA /GEN=LRRFIP2 /PROD=leucine rich repeat (in FLII) interacting protein 2 /DB_XREF=gi:8923223 /UG=Hs.57672 leucine rich repeat (in FLII) interacting protein 2 /FL=gb:NM_017724.1	NM_017724		NP_060194
218375_at	0.033533	gb:NM_024047.1 /DEF=Homo sapiens hypothetical protein MGC3037 (MGC3037), mRNA. /FEA=mRNA /GEN=MGC3037 /PROD=hypothetical protein MGC3037 /DB_XREF=gi:13129009 /UG=Hs.301789 hypothetical protein MGC3037 /FL=gb:BC000542.1 gb:NM_024047.1	NM_024047		NP_076952
218376_s_at	0.013223	gb:NM_022765.1 /DEF=Homo sapiens hypothetical protein FLJ11937 (FLJ11937), mRNA. /FEA=mRNA /GEN=FLJ11937 /PROD=hypothetical protein FLJ11937 /DB_XREF=gi:12232438 /UG=Hs.33476 hypothetical protein FLJ11937 /FL=gb:NM_022765.1	NM_022765		NP_073602
218379_at	0.007779	gb:NM_016090.1 /DEF=Homo sapiens RNA binding motif protein 7 (RBM7), mRNA. /FEA=mRNA /GEN=RBM7 /PROD=RNA binding motif protein 7 /DB_XREF=gi:9994184 /UG=Hs.5887 RNA binding motif protein 7 /FL=gb:AF156098.1 gb:NM_016090.1	NM_016090		NP_057174

G n Id ntifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
218383_at	0.001116	gb:NM_017815.1 /DEF=Homo sapiens hypothetical protein FLJ20424 (FLJ20424), mRNA. /FEA=mRNA /GEN=FLJ20424 /PROD=hypothetical protein FLJ20424 /DB_XREF=gi:8923395 /UG=Hs.8886 hypothetical protein FLJ20424 /FL=gb:BC002554.1 gb:BC001916.1 gb:NM_017815.1	NM_017815		NP_060285
218394_at	0.00161	gb:NM_024589.1 /DEF=Homo sapiens hypothetical protein FLJ22386 (FLJ22386), mRNA. /FEA=mRNA /GEN=FLJ22386 /PROD=hypothetical protein FLJ22386 /DB_XREF=gi:13375778 /UG=Hs.22795 hypothetical protein FLJ22386 /FL=gb:AL136675.1 gb:NM_024589.1	NM_024589		NP_078865
218399_s_at	0.033793	gb:NM_017955.1 /DEF=Homo sapiens hypothetical protein FLJ20764 (FLJ20764), mRNA. /FEA=mRNA /GEN=FLJ20764 /PROD=hypothetical protein FLJ20764 /DB_XREF=gi:8923673 /UG=Hs.34045 hypothetical protein FLJ20764 /FL=gb:NM_017955.1	NM_017955		NP_663747
218404_at	0.001116	gb:NM_013322.1 /DEF=Homo sapiens sorting nexin 10 (SNX10), mRNA. /FEA=mRNA /GEN=SNX10 /PROD=sorting nexin 10 /DB_XREF=gi:7019536 /UG=Hs.106260 sorting nexin 10 /FL=gb:AF121860.1 gb:NM_013322.1	NM_013322		NP_037454
218414_s_at	5.03E-04	gb:NM_017668.1 /DEF=Homo sapiens LIS1-interacting protein NUDE1, rat homolog (FLJ20101), mRNA. /FEA=mRNA /GEN=FLJ20101 /PROD=LIS1-interacting protein NUDE1, rat homolog /DB_XREF=gi:8923109 /UG=Hs.263925 LIS1-interacting protein NUDE1, rat homolog /FL=gb:BC001421.1 gb:NM_017668.1	NM_017668		NP_060138

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218419_s_at	0.02145	gb:NM_024107.1 /DEF=Homo sapiens hypothetical protein MGC3123 (MGC3123), mRNA. /FEA=mRNA /GEN=MGC3123 /PROD=hypothetical protein MGC3123 /DB_XREF=gi:13129117 /UG=Hs.288600 hypothetical protein MGC3123 /FL=gb:BC001792.1 gb:NM_024107.1	NM_024107		NP_803190
218420_s_at	0.037567	gb:NM_025138.1 /DEF=Homo sapiens hypothetical protein FLJ12661 (FLJ12661), mRNA. /FEA=mRNA /GEN=FLJ12661 /PROD=hypothetical protein FLJ12661 /DB_XREF=gi:13376720 /UG=Hs.318526 hypothetical protein FLJ12661 /FL=gb:NM_025138.1	NM_025138		NP_733837
218423_x_at	1.26E-04	gb:NM_016516.1 /DEF=Homo sapiens tumor antigen SLP-8p (HCC8), mRNA. /FEA=mRNA /GEN=HCC8 /PROD=tumor antigen SLP-8p /DB_XREF=gi:7705396 /UG=Hs.48499 tumor antigen SLP-8p /FL=gb:AF102177.1 gb:NM_016516.1	NM_016516		NP_057600
218424_s_at	0.008237	gb:NM_018234.1 /DEF=Homo sapiens hypothetical protein FLJ10829 (FLJ10829), mRNA. /FEA=mRNA /GEN=FLJ10829 /PROD=hypothetical protein FLJ10829 /DB_XREF=gi:8922696 /UG=Hs.57655 hypothetical protein FLJ10829 /FL=gb:NM_018234.1	NM_018234		NP_060704
218426_s_at	0.00161	gb:NM_019011.1 /DEF=Homo sapiens TRIAD3 protein (TRIAD3), mRNA. /FEA=mRNA /GEN=TRIAD3 /PROD=TRIAD3 protein /DB_XREF=gi:9507204 /UG=Hs.86228 TRIAD3 protein /FL=gb:BC000787.1 gb:NM_019011.1	NM_019011		NP_061884

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218431_at	0.02145	gb:NM_022067.1 /DEF=Homo sapiens hypothetical protein FLJ12707 (FLJ12707), mRNA. /FEA=mRNA /GEN=FLJ12707 /PROD=hypothetical protein FLJ12707 /DB_XREF=gi:11545778 /UG=Hs.16157 hypothetical protein FLJ12707 /FL=gb:NM_022067.1	NM_022067		NP_071350
218439_s_at	0.003817	gb:NM_016144.1 /DEF=Homo sapiens PTD002 protein (PTD002), mRNA. /FEA=mRNA /GEN=PTD002 /PROD=PTD002 protein /DB_XREF=gi:7706662 /UG=Hs.41767 PTD002 protein /FL=gb:BC005179.1 gb:AF078857.1 gb:NM_016144.1	NM_016144		NP_057228
218448_at	0.010205	gb:NM_017896.1 /DEF=Homo sapiens hypothetical protein FLJ20602 (FLJ20602), mRNA. /FEA=mRNA /GEN=FLJ20602 /PROD=hypothetical protein FLJ20602 /DB_XREF=gi:8923556 /UG=Hs.103808 hypothetical protein FLJ20602 /FL=gb:NM_017896.1	NM_017896		NP_060366
218450_at	0.004341	gb:NM_015987.1 /DEF=Homo sapiens heme-binding protein (HEBP), mRNA. /FEA=mRNA /GEN=HEBP /PROD=heme-binding protein /DB_XREF=gi:7705404 /UG=Hs.108675 heme-binding protein /FL=gb:AF117615.1 gb:NM_015987.1 gb:AF167473.1	NM_015987		NP_057071
218454_at	1.26E-04	gb:NM_024829.1 /DEF=Homo sapiens hypothetical protein FLJ22662 (FLJ22662), mRNA. /FEA=mRNA /GEN=FLJ22662 /PROD=hypothetical protein FLJ22662 /DB_XREF=gi:13376231 /UG=Hs.178470 hypothetical protein FLJ22662 /FL=gb:NM_024829.1	NM_024829		NP_079105

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
218458_at	1.26E-04	gb:NM_022471.1 /DEF=Homo sapiens hypothetical protein FLJ13057 similar to germ cell-less (FLJ13057), mRNA. /FEA=mRNA /GEN=FLJ13057 /PROD=hypothetical protein FLJ13057 similar to germcell-less /DB_XREF=gi:11968020 /UG=Hs.243122 hypothetical protein FLJ13057 similar to germ cell-less /FL=gb:NM_022471.1	NM_022471		NP_848526
218465_at	2.05E-04	gb:NM_018126.1 /DEF=Homo sapiens hypothetical protein FLJ10525 (FLJ10525), mRNA. /FEA=mRNA /GEN=FLJ10525 /PROD=hypothetical protein FLJ10525 /DB_XREF=gi:8922490 /UG=Hs.31082 hypothetical protein FLJ10525 /FL=gb:BC000948.1 gb:NM_018126.1	NM_018126		NP_060596
218471_s_at	0.001886	gb:NM_024649.1 /DEF=Homo sapiens hypothetical protein FLJ23590 (FLJ23590), mRNA. /FEA=mRNA /GEN=FLJ23590 /PROD=hypothetical protein FLJ23590 /DB_XREF=gi:13375892 /UG=Hs.54890 hypothetical protein FLJ23590 /FL=gb:NM_024649.1	NM_024649		NP_078925
218472_s_at	5.03E-04	gb:NM_015946.1 /DEF=Homo sapiens pelota (Drosophila) homolog (PELO), mRNA. /FEA=mRNA /GEN=PELO /PROD=CGI-17 protein /DB_XREF=gi:7705599 /UG=Hs.5798 pelota (Drosophila) homolog /FL=gb:AF139828.1 gb:AF132951.1 gb:NM_015946.1	NM_015946		NP_057030
218482_at	2.05E-04	gb:NM_020189.1 /DEF=Homo sapiens DC6 protein (DC6), mRNA. /FEA=mRNA /GEN=DC6 /PROD=DC6 protein /DB_XREF=gi:9910185 /UG=Hs.283740 DC6 protein /FL=gb:AF201940.1 gb:AF173296.1 gb:NM_020189.1	NM_020189		NP_064574

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218487_at	2.05E-04	gb:BC000977.1 /DEF=Homo sapiens, aminolevulinate, delta-, dehydratase, clone MGC:5057, mRNA, complete cds. /FEA=mRNA /PROD=aminolevulinate, delta-, dehydratase /DB_XREF=gi:12654312 /UG=Hs.1227 aminolevulinate, delta-, dehydratase /FL=gb:BC000977.1 gb:M13928.1 gb:NM_000031.1	BC000977		NP_000022
218490_s_at	0.022795	gb:NM_018443.1 /DEF=Homo sapiens zinc finger protein ZNF140-like protein (LOC55828), mRNA. /FEA=mRNA /GEN=LOC55828 /PROD=zinc finger protein ZNF140-like protein /DB_XREF=gi:8923850 /UG=Hs.125287 zinc finger protein ZNF140-like protein /FL=gb:AF155656.1 gb:NM_018443.1	NM_018443		NP_061145
218492_s_at	0.020793	gb:NM_030573.1 /DEF=Homo sapiens hypothetical protein MGC10963 (MGC10963), mRNA. /FEA=mRNA /GEN=MGC10963 /PROD=hypothetical protein MGC10963 /DB_XREF=gi:13386483 /UG=Hs.14927 hypothetical protein MGC10963. /FL=gb:BC004346.1 gb:NM_030573.1	NM_030573		NP_085050
218493_at	0.025958	gb:NM_024571.1 /DEF=Homo sapiens hypothetical protein FLJ22940 (FLJ22940), mRNA. /FEA=mRNA /GEN=FLJ22940 /PROD=hypothetical protein FLJ22940 /DB_XREF=gi:13443017 /UG=Hs.15277 hypothetical protein FLJ22940 /FL=gb:BC001381.1 gb:NM_024571.1	NM_024571		NP_078847

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218498_s_at	1.32E-04	gb:NM_014584.1 /DEF=Homo sapiens ERO1 (S. cerevisiae)-like (ERO1L), mRNA. /FEA=mRNA /GEN=ERO1L /PROD=ERO1 (S. cerevisiae)-like /DB_XREF=gi:7657068 /UG=Hs.25740 ERO1 (S. cerevisiae)-like /FL=gb:AF081886.1 gb:NM_014584.1	NM_014584		NP_055399
218499_at	0.026891	gb:NM_016542.1 /DEF=Homo sapiens serinethreonine protein kinase MASK (LOC51765), mRNA. /FEA=mRNA /GEN=LOC51765 /PROD=serinethreonine protein kinase MASK /DB_XREF=gi:7706568 /UG=Hs.23643 serinethreonine protein kinase MASK /FL=gb:AB040057.1 gb:NM_016542.1	NM_016542		NP_057626
218502_s_at	0.005852	gb:NM_014112.1 /DEF=Homo sapiens trichorhinophalangeal syndrome I gene (TRPS1), mRNA. /FEA=mRNA /GEN=TRPS1 /PROD=trichorhinophalangeal syndrome I gene /DB_XREF=gi:7657658 /UG=Hs.26102 trichorhinophalangeal syndrome I /FL=gb:AF183810.1 gb:NM_014112.1	NM_014112		NP_054831
218506_x_at	2.05E-04	gb:NM_018459.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM045 (BM045), mRNA. /FEA=mRNA /GEN=BM045 /PROD=uncharacterized bone marrow protein BM045 /DB_XREF=gi:8922103 /UG=Hs.8750 uncharacterized bone marrow protein BM045 /FL=gb:AF217521.1 gb:NM_018459.1	NM_018459		NP_060929

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218508_at	5.64E-04	gb:NM_018403.1 /DEF=Homo sapiens transcription factor (SMIF gene), (HSA275986), mRNA. /FEA=mRNA /GEN=HSA275986 /PROD=transcription factor (SMIF gene) /DB_XREF=gi:8923766 /UG=Hs.71414 transcription factor (SMIF gene) /FL=gb:NM_018403.1	NM_018403		NP_060873
218514_at	3.26E-04	gb:NM_018149.1 /DEF=Homo sapiens hypothetical protein FLJ10587 (FLJ10587), mRNA. /FEA=mRNA /GEN=FLJ10587 /PROD=hypothetical protein FLJ10587 /DB_XREF=gi:8922539 /UG=Hs.7296 hypothetical protein FLJ10587 /FL=gb:NM_018149.1	NM_018149		NP_060619
218518_at	0.016934	gb:NM_016603.1 /DEF=Homo sapiens GAP-like protein (LOC51306), mRNA. /FEA=mRNA /GEN=LOC51306 /PROD=GAP-like protein /DB_XREF=gi:7706136 /UG=Hs.82035 potential nuclear protein C5ORF5; GAP-like protein /FL=gb:AF251038.1 gb:AF157316.1 gb:NM_016603.1	NM_016603		NP_057687
218519_at	0.005852	gb:NM_017945.1 /DEF=Homo sapiens hypothetical protein FLJ20730 (FLJ20730), mRNA. /FEA=mRNA /GEN=FLJ20730 /PROD=hypothetical protein FLJ20730 /DB_XREF=gi:8923656 /UG=Hs.237480 hypothetical protein FLJ20730 /FL=gb:BC005207.1 gb:NM_017945.1	NM_017945		NP_060415
218520_at	0.005852	gb:NM_013254.1 /DEF=Homo sapiens TANK-binding kinase 1 (TBK1), mRNA. /FEA=mRNA /GEN=TBK1 /PROD=TANK-binding kinase 1 /DB_XREF=gi:7019546 /UG=Hs.21712 TANK-binding kinase 1 /FL=gb:AF191838.1 gb:NM_013254.1 gb:AF174536.1	NM_013254		NP_037386

Gen Id ntifier	p-value	Description	Gene Accession No.	Unigene Acc ssion No.	Protein Accession No.
218522_s_at	0.001116	gb:NM_018174.1 /DEF=Homo sapiens hypothetical protein FLJ10669 (FLJ10669), mRNA. /FEA=mRNA /GEN=FLJ10669 /PROD=hypothetical protein FLJ10669 /DB_XREF=gi:8922582 /UG=Hs.66048 hypothetical protein FLJ10669 /FL=gb:NM_018174.1	NM_018174		NP_060644
218526_s_at	0.029453	gb:NM_014185.1 /DEF=Homo sapiens HSPC165 protein (HSPC165), mRNA. /FEA=mRNA /GEN=HSPC165 /PROD=HSPC165 protein /DB_XREF=gi:7661825 /UG=Hs.13605 HSPC165 protein /FL=gb:AF161514.1 gb:AF151070.1 gb:NM_014185.1 gb:NM_016492.1 gb:AF168714.1 gb:AF265206.1	NM_014185		NP_057576
218528_s_at	1.26E-04	gb:NM_022781.1 /DEF=Homo sapiens hypothetical protein FLJ21343 (FLJ21343), mRNA. /FEA=mRNA /GEN=FLJ21343 /PROD=hypothetical protein FLJ21343 /DB_XREF=gi:12232470 /UG=Hs.77823 hypothetical protein FLJ21343 /FL=gb:AL136817.1 gb:NM_022781.1	NM_022781		NP_073618
218531_at	0.013223	gb:NM_025124.1 /DEF=Homo sapiens hypothetical protein FLJ21749 (FLJ21749), mRNA. /FEA=mRNA /GEN=FLJ21749 /PROD=hypothetical protein FLJ21749 /DB_XREF=gi:13376700 /UG=Hs.288761 hypothetical protein FLJ21749 /FL=gb:NM_025124.1	NM_025124		NP_079400
218534_s_at	7.58E-04	gb:NM_018046.1 /DEF=Homo sapiens hypothetical protein FLJ10283 (FLJ10283), mRNA. /FEA=mRNA /GEN=FLJ10283 /PROD=hypothetical protein FLJ10283 /DB_XREF=gi:8922325 /UG=Hs.284216 hypothetical protein FLJ10283 /FL=gb:NM_018046.1	NM_018046		NP_060516

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218539_at	0.026891	gb:NM_017943.1 /DEF=Homo sapiens hypothetical protein FLJ20725 (FLJ20725), mRNA. /FEA=mRNA /GEN=FLJ20725 /PROD=hypothetical protein FLJ20725 /DB_XREF=gi:8923650 /UG=Hs.15467 hypothetical protein. FLJ20725 /FL=gb:NM_017943.1	NM_017943		NP_060413
218549_s_at	1.26E-04	gb:NM_016033.1 /DEF=Homo sapiens CGI-90 protein (LOC51115), mRNA. /FEA=mRNA /GEN=LOC51115 /PROD=CGI-90 protein /DB_XREF=gi:7705802 /UG=Hs.44222 CGI-90 protein /FL=gb:AF151848.1 gb:NM_016033.1	NM_016033		NP_057117
218556_at	1.26E-04	gb:NM_014182.1 /DEF=Homo sapiens HSPC160 protein (HSPC160), mRNA. /FEA=mRNA /GEN=HSPC160 /PROD=HSPC160 protein /DB_XREF=gi:7661819 /UG=Hs.13144 HSPC160 protein /FL=gb:AF161509.1 gb:NM_014182.1	NM_014182		NP_054901
218563_at	0.013223	gb:NM_004542.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9kD, B9) (NDUFA3), mRNA. /FEA=mRNA /GEN=NDUFA3 /PROD=NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9kD, B9) /DB_XREF=gi:4758771 /UG=Hs.198269 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9kD, B9) /FL=gb:AF044955.1 gb:AF070653.1 gb:NM_004542.1	NM_004542		NP_004533
218572_at	0.021435	gb:NM_014169.1 /DEF=Homo sapiens HSPC134 protein (HSPC134), mRNA. /FEA=mRNA /GEN=HSPC134 /PROD=HSPC134 protein /DB_XREF=gi:7661793 /UG=Hs.279761 HSPC134 protein /FL=gb:AF212243.1 gb:AF161483.1 gb:NM_014169.1	NM_014169		NP_054888

G n Identifier	p-valu	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218578_at	1.32E-04	gb:NM_024529.1 /DEF=Homo sapiens hypothetical protein FLJ23316 (FLJ23316), mRNA. /FEA=mRNA /GEN=FLJ23316 /PROD=hypothetical protein FLJ23316 /DB_XREF=gi:13375677 /UG=Hs.5722 hypothetical protein FLJ23316 /FL=gb:AF312865.1 gb:NM_024529.1	NM_024529		NP_078805
218580_x_at	0.007779	gb:NM_017900.1 /DEF=Homo sapiens hypothetical protein FLJ20608 (FLJ20608), mRNA. /FEA=mRNA /GEN=FLJ20608 /PROD=hypothetical protein FLJ20608 /DB_XREF=gi:8923564 /UG=Hs.76239 hypothetical protein FLJ20608 /FL=gb:NM_017900.1	NM_017900		NP_060370
218582_at	2.05E-04	gb:NM_017824.1 /DEF=Homo sapiens hypothetical protein FLJ20445 (FLJ20445), mRNA. /FEA=mRNA /GEN=FLJ20445 /PROD=hypothetical protein FLJ20445 /DB_XREF=gi:8923414 /UG=Hs.10340 hypothetical protein FLJ20445 /FL=gb:NM_017824.1	NM_017824		NP_060294
218583_s_at	2.05E-04	gb:NM_020640.1 /DEF=Homo sapiens RP42 homolog (RP42), mRNA. /FEA=mRNA /GEN=RP42 /PROD=RP42 homolog /DB_XREF=gi:10190677 /UG=Hs.104613 RP42 homolog /FL=gb:NM_020640.1 gb:AF292100.2	NM_020640		NP_065691
218586_at	0.010205	gb:NM_018270.1 /DEF=Homo sapiens hypothetical protein FLJ10914 (FLJ10914), mRNA. /FEA=mRNA /GEN=FLJ10914 /PROD=hypothetical protein FLJ10914 /DB_XREF=gi:8922763 /UG=Hs.143954 hypothetical protein FLJ10914 /FL=gb:NM_018270.1	NM_018270		NP_060740

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218588_s_at	0.00161	gb:NM_018691.1 /DEF=Homo sapiens chromosome 5 open reading frame 3 (C5ORF3), mRNA. /FEA=mRNA /GEN=C5ORF3 /PROD=hypothetical protein /DB_XREF=gi:8922068 /UG=Hs.166551 chromosome 5 open reading frame 3 /FL=gb:NM_018691.1	NM_018691		NP_061161
218597_s_at	0.001477	gb:NM_018464.1 /DEF=Homo sapiens uncharacterized hematopoietic stemprogenitor cells protein MDS029 (MDS029), mRNA. /FEA=mRNA /GEN=MDS029 /PROD=uncharacterized hematopoietic stemprogenitorcells protein MDS029 /DB_XREF=gi:8923929 /UG=Hs.43549 uncharacterized hematopoietic stemprogenitor cells protein MDS029 /FL=gb:AF220049.1 gb:NM_018464.1	NM_018464		NP_060934
218603_at	0.001116	gb:NM_016217.1 /DEF=Homo sapiens hHDC for homolog of Drosophila headcase (LOC51696), mRNA. /FEA=mRNA /GEN=LOC51696 /PROD=hHDC for homolog of Drosophila headcase /DB_XREF=gi:7706434 /UG=Hs.6679 hHDC for homolog of Drosophila headcase /FL=gb:AB033492.1 gb:NM_016217.1	NM_016217		NP_057301
218606_at	1.26E-04	gb:NM_017740.1 /DEF=Homo sapiens hypothetical protein FLJ20279. (FLJ20279), mRNA. /FEA=mRNA /GEN=FLJ20279 /PROD=hypothetical protein FLJ20279 /DB_XREF=gi:8923254 /UG=Hs.9725 hypothetical protein FLJ20279 /FL=gb:NM_017740.1	NM_017740		NP_060210

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
218616_at	2.05E-04	gb:NM_020395.1 /DEF=Homo sapiens hypothetical nuclear factor SBBI22 (LOC57117), mRNA. /FEA=mRNA /GEN=LOC57117 /PROD=hypothetical nuclear factor SBBI22 /DB_XREF=gi:9966872 /UG=Hs.26323 hypothetical nuclear factor SBBI22 /FL=gb:AF242524.1 gb:NM_020395.1	NM_020395		NP_065128
218617_at	0.035611	gb:NM_017646.1 /DEF=Homo sapiens tRNA isopentenylpyrophosphate transferase (IPT), mRNA. /FEA=mRNA /GEN=IPT /PROD=tRNA isopentenylpyrophosphate transferase. /DB_XREF=gi:8923064 /UG=Hs.288036 tRNA isopentenylpyrophosphate transferase /FL=gb:NM_017646.1	NM_017646		NP_060116
218625_at	0.002864	gb:NM_016588.1 /DEF=Homo sapiens neuritin (LOC51299), mRNA. /FEA=mRNA /GEN=LOC51299 /PROD=neuritin /DB_XREF=gi:7706122 /UG=Hs.103291 neuritin /FL=gb:BC002683.1 gb:AF136631.1 gb:NM_016588.1	NM_016588		NP_057672
218627_at	1.26E-04	gb:NM_018370.1 /DEF=Homo sapiens hypothetical protein FLJ11259 (FLJ11259), mRNA. /FEA=mRNA /GEN=FLJ11259 /PROD=hypothetical protein FLJ11259 /DB_XREF=gi:8922957 /UG=Hs.184465 hypothetical protein FLJ11259 /FL=gb:NM_018370.1	NM_018370		NP_060840

Gene Identifier	p-value	Description	Gene Accession No.	Unig n Acc ssion No.	Protein Accession No.
218628_at	0.001116	gb:NM_016053.1 /DEF=Homo sapiens CGI-116 protein (LOC51019), mRNA. /FEA=mRNA /GEN=LOC51019 /PROD=CGI-116 protein /DB_XREF=gi:7705621 /UG=Hs.18885 CGI-116 protein /FL=gb:AF151874.1 gb:NM_016053.1	NM_016053		NP_057137
218632_at	0.026891	gb:NM_024602.1 /DEF=Homo sapiens hypothetical protein FLJ21156 (FLJ21156), mRNA. /FEA=mRNA /GEN=FLJ21156 /PROD=hypothetical protein FLJ21156 /DB_XREF=gi:13375804 /UG=Hs.26058 hypothetical protein FLJ21156 /FL=gb:NM_024602.1	NM_024602		NP_078878
218637_at	0.033381	gb:NM_018439.1 /DEF=Homo sapiens hypothetical protein IMPACT (IMPACT), mRNA. /FEA=mRNA /GEN=IMPACT /PROD=hypothetical protein IMPACT /DB_XREF=gi:8923818 /UG=Hs.284245 hypothetical protein IMPACT /FL=gb:AF208694.1 gb:AB026264.1 gb:NM_018439.1	NM_018439		NP_060909
218639_s_at	0.035171	gb:NM_025112.1 /DEF=Homo sapiens hypothetical protein MGC11349 (MGC11349), mRNA. /FEA=mRNA /GEN=MGC11349 /PROD=hypothetical protein MGC11349 /DB_XREF=gi:13376684 /UG=Hs.288697 hypothetical protein MGC11349 /FL=gb:BC002940.1 gb:NM_025112.1	NM_025112		NP_079388

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218645_at	0.046906	gb:NM_021994.1 /DEF=Homo sapiens zinc finger protein 277 (ZNF277), mRNA. /FEA=mRNA /GEN=ZNF277 /PROD=zinc finger protein 277 /DB_XREF=gi:11496268 /UG=Hs.42636 zinc finger protein 277 /FL=gb:NM_021994.1 gb:AF209198.1	NM_021994		NP_068834
218652_s_at	0.007779	gb:NM_017733.1 /DEF=Homo sapiens hypothetical protein FLJ20265 (FLJ20265), mRNA. /FEA=mRNA /GEN=FLJ20265 /PROD=hypothetical protein FLJ20265 /DB_XREF=gi:8923239 /UG=Hs.7099 hypothetical protein FLJ20265 /FL=gb:BC001249.1 gb:BC000937.2 gb:NM_017733.1	NM_017733		NP_060203
218660_at	3.26E-04	gb:NM_003494.1 /DEF=Homo sapiens dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive) (DYSF), mRNA. /FEA=mRNA /GEN=DYSF /PROD=dysferlin /DB_XREF=gi:4503430 /UG=Hs.143897 dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive) /FL=gb:AF075575.1 gb:NM_003494.1	NM_003494		NP_003485
218661_at	5.64E-04	gb:NM_024845.1 /DEF=Homo sapiens hypothetical protein FLJ14154 (FLJ14154), mRNA. /FEA=mRNA /GEN=FLJ14154 /PROD=hypothetical protein FLJ14154 /DB_XREF=gi:13376260 /UG=Hs.184870 hypothetical protein FLJ14154 /FL=gb:NM_024845.1	NM_024845		NP_079121

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218662_s_at	2.05E-04	gb:NM_022346.1 /DEF=Homo sapiens chromosome condensation protein G (HCAP-G), mRNA. /FEA=mRNA /GEN=HCAP-G /PROD=chromosome condensation protein G /DB_XREF=gi:11641252 /UG=Hs.193602 chromosome condensation protein G /FL=gb:AF235023.1 gb:NM_022346.1 gb:AF331796.1 gb:BC000827.1 gb:AB013299.1	NM_022346		NP_071741
218668_s_at	3.26E-04	gb:NM_021183.1 /DEF=Homo sapiens hypothetical protein similar to small G proteins, especially RAP-2A (LOC57826), mRNA. /FEA=mRNA /GEN=LOC57826 /PROD=hypothetical protein similar to small Gproteins, especially RAP-2A /DB_XREF=gi:10880976 /UG=Hs.225979 hypothetical protein similar to small G proteins, especially RAP-2A /FL=gb:NM_021183.1 gb:BC003403.1	NM_021183		NP_067006
218669_at	0.00161	gb:NM_021183.1 /DEF=Homo sapiens hypothetical protein similar to small G proteins, especially RAP-2A (LOC57826), mRNA. /FEA=mRNA /GEN=LOC57826 /PROD=hypothetical protein similar to small Gproteins, especially RAP-2A /DB_XREF=gi:10880976 /UG=Hs.225979 hypothetical protein similar to small G proteins, especially RAP-2A /FL=gb:NM_021183.1 gb:BC003403.1	NM_021183		NP_067006

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Prot in Accession No.
218673_s_at	0.041055	gb:NM_006395.1 /DEF=Homo sapiens ubiquitin activating enzyme E1-like protein (GSA7), mRNA. /FEA=mRNA /GEN=GSA7 /PROD=ubiquitin activating enzyme E1-like protein /DB_XREF=gi:5453667 /UG=Hs.278607 ubiquitin activating enzyme E1-like protein /FL=gb:AF094516.1 gb:NM_006395.1	NM_006395		NP_006386
218676_s_at	0.005852	gb:NM_021213.1 /DEF=Homo sapiens phosphatidylcholine transfer protein (PCTP), mRNA. /FEA=mRNA /GEN=PCTP /PROD=phosphatidylcholine transfer protein /DB_XREF=gi:10864026 /UG=Hs.285218 phosphatidylcholine transfer protein /FL=gb:NM_021213.1 gb:AF151638.1	NM_021213		NP_067036
218679_s_at	0.004341	gb:NM_016208.1 /DEF=Homo sapiens VPS28 protein (LOC51160), mRNA. /FEA=mRNA /GEN=LOC51160 /PROD=VPS28 protein /DB_XREF=gi:7705884 /UG=Hs.293441 VPS28 protein /FL=gb:AF316887.1 gb:AF182844.1 gb:NM_016208.1	NM_016208		NP_057292
218682_s_at	1.26E-04	gb:NM_018158.1 /DEF=Homo sapiens hypothetical protein FLJ10624 (FLJ10624), mRNA. /FEA=mRNA /GEN=FLJ10624 /PROD=hypothetical protein FLJ10624 /DB_XREF=gi:8922556 /UG=Hs.306000 hypothetical protein FLJ10624 /FL=gb:NM_018158.1	NM_018158		NP_060628
218698_at	0.02145	gb:NM_015957.1 /DEF=Homo sapiens CGI-29 protein (LOC51074), mRNA. /FEA=mRNA /GEN=LOC51074 /PROD=CGI-29 protein /DB_XREF=gi:7705723 /UG=Hs.104058 CGI-29 protein /FL=gb:AF132963.1 gb:NM_015957.1	NM_015957		NP_057041

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218703_at	0.021448	gb:NM_012430.1 /DEF=Homo sapiens sec22 homolog (SEC22A), mRNA. /FEA=mRNA /GEN=SEC22A /PROD=sec22 homolog /DB_XREF=gi:6912647 /UG=Hs.183655 sec22 homolog /FL=gb:AF100749.1 gb:NM_012430.1	NM_012430		NP_036562
218718_at	3.26E-04	gb:NM_016205.1 /DEF=Homo sapiens platelet derived growth factor C (PDGFC), mRNA. /FEA=mRNA /GEN=PDGFC /PROD=secretory growth factor-like protein fallotein /DB_XREF=gi:9994186 /UG=Hs.43080 platelet derived growth factor C. /FL=gb:AF091434.1 gb:AF244813.1 gb:AB033831.1 gb:NM_016205.1	NM_016205		NP_057289
218722_s_at	0.025245	gb:NM_024661.1 /DEF=Homo sapiens hypothetical protein FLJ12436 (FLJ12436), mRNA. /FEA=mRNA /GEN=FLJ12436 /PROD=hypothetical protein FLJ12436 /DB_XREF=gi:13375914 /UG=Hs.69485 hypothetical protein FLJ12436 /FL=gb:NM_024661.1	NM_024661		NP_078937
218723_s_at	0.016934	gb:NM_014059.1 /DEF=Homo sapiens RGC32 protein (RGC32), mRNA. /FEA=mRNA /GEN=RGC32 /PROD=RGC32 protein /DB_XREF=gi:7662650 /UG=Hs.76640 RGC32 protein /FL=gb:AF036549.1 gb:NM_014059.1	NM_014059		NP_054778
218724_s_at	0.003539	gb:NM_021809.1 /DEF=Homo sapiens TGF(beta)-induced transcription factor 2 (TGIF2), mRNA. /FEA=mRNA /GEN=TGIF2 /PROD=TGF(beta)-induced transcription factor 2 /DB_XREF=gi:11141880 /UG=Hs.94785 TGF(beta)-induced transcription factor 2 /FL=gb:AB042646.1 gb:NM_021809.1	NM_021809		NP_068581

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
218728_s_at	1.26E-04	gb:NM_014184.1 /DEF=Homo sapiens HSPC163 protein (HSPC163), mRNA. /FEA=mRNA /GEN=HSPC163 /PROD=HSPC163 protein /DB_XREF=gi:7661823 /UG=Hs.108854 HSPC163 protein /FL=gb:BC000573.1 gb:AF161512.1 gb:NM_014184.1	NM_014184		NP_054903
218729_at	0.007258	gb:NM_020169.1 /DEF=Homo sapiens latexin protein (LXN), mRNA. /FEA=mRNA /GEN=LXN /PROD=latexin protein /DB_XREF=gi:9910395 /UG=Hs.109276 latexin protein /FL=gb:BC005346.1 gb:AF282626.1 gb:NM_020169.1	NM_020169		NP_064554
218734_at	0.004341	gb:NM_024771.1 /DEF=Homo sapiens hypothetical protein FLJ13848 (FLJ13848), mRNA. /FEA=mRNA /GEN=FLJ13848 /PROD=hypothetical protein FLJ13848 /DB_XREF=gi:13376118 /UG=Hs.136976 hypothetical protein FLJ13848 /FL=gb:NM_024771.1	NM_024771		NP_079047
218738_s_at	5.03E-04	gb:NM_016271.1 /DEF=Homo sapiens STRIN protein (STRIN), mRNA. /FEA=mRNA /GEN=STRIN /PROD=STRIN protein /DB_XREF=gi:7706722 /UG=Hs.180403 STRIN protein /FL=gb:AF162680.3 gb:NM_016271.1	NM_016271		NP_057355
218739_at	2.05E-04	gb:NM_016006.1 /DEF=Homo sapiens CGI-58 protein (LOC51099), mRNA. /FEA=mRNA /GEN=LOC51099 /PROD=CGI-58 protein /DB_XREF=gi:7705770 /UG=Hs.19385 CGI-58 protein /FL=gb:AF151816.1 gb:NM_016006.1	NM_016006		NP_057090

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218761_at	7.58E-04	gb:NM_017610.1 /DEF=Homo sapiens hypothetical protein DKFZp761D081 (DKFZp761D081), mRNA. /FEA=mRNA /GEN=DKFZp761D081 /PROD=hypothetical protein DKFZp761D081 /DB_XREF=gi:8922164 /UG=Hs.12504 hypothetical protein DKFZp761D081 /FL=gb:NM_017610.1	NM_017610		NP_060080
218764_at	5.03E-04	gb:NM_024064.1 /DEF=Homo sapiens hypothetical protein MGC5363 (MGC5363), mRNA. /FEA=mRNA /GEN=MGC5363 /PROD=hypothetical protein MGC5363 /DB_XREF=gi:13129041 /UG=Hs.1880 hypothetical protein MGC5363 /FL=gb:BC001000.2 gb:NM_024064.1	NM_024064		
218773_s_at	7.58E-04	gb:NM_012228.1 /DEF=Homo sapiens pilin-like transcription factor (PILB), mRNA. /FEA=mRNA /GEN=PILB /PROD=pilin-like transcription factor /DB_XREF=gi:6912591 /UG=Hs.279754 pilin-like transcription factor /FL=gb:AF122004.1 gb:NM_012228.1	NM_012228		NP_036360
218782_s_at	0.013223	gb:NM_014109.1 /DEF=Homo sapiens PRO2000 protein (PRO2000), mRNA. /FEA=mRNA /GEN=PRO2000 /PROD=PRO2000 protein /DB_XREF=gi:7662630 /UG=Hs.46677 PRO2000 protein /FL=gb:AF118088.1 gb:NM_014109.1	NM_014109		NP_054828
218789_s_at	0.001848	gb:NM_019021.1 /DEF=Homo sapiens hypothetical protein (FLJ20010), mRNA. /FEA=mRNA /GEN=FLJ20010 /PROD=hypothetical protein /DB_XREF=gi:9506646 /UG=Hs.91816 hypothetical protein /FL=gb:NM_019021.1	NM_019021		NP_061894

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218799_at	0.001205	gb:NM_018066.1 /DEF=Homo sapiens hypothetical protein FLJ10349 (FLJ10349), mRNA. /FEA=mRNA /GEN=FLJ10349 /PROD=hypothetical protein FLJ10349 /DB_XREF=gi:8922365 /UG=Hs.19333 hypothetical protein FLJ10349 /FL=gb:NM_018066.1	NM_018066		NP_060536
218805_at	0.001116	gb:NM_018384.1 /DEF=Homo sapiens hypothetical protein FLJ11296 (FLJ11296), mRNA. /FEA=mRNA /GEN=FLJ11296 /PROD=hypothetical protein FLJ11296 /DB_XREF=gi:8922984 /UG=Hs.26194 hypothetical protein FLJ11296 /FL=gb:NM_018384.1	NM_018384		NP_060854
218806_s_at	2.32E-04	gb:AF118887.1 /DEF=Homo sapiens VAV-3 protein (VAV-3) mRNA, alternatively spliced, complete cds. /FEA=mRNA /GEN=VAV-3 /PROD=VAV-3 protein /DB_XREF=gi:4416407 /UG=Hs.267659 vav 3 oncogene /FL=gb:AF067817.1 gb:AF118887.1 gb:NM_006113.2	AF118887		NP_006104
218809_at	0.005852	gb:NM_024960.1 /DEF=Homo sapiens hypothetical protein FLJ11729 (FLJ11729), mRNA. /FEA=mRNA /GEN=FLJ11729 /PROD=hypothetical protein FLJ11729 /DB_XREF=gi:13376450 /UG=Hs.286212 hypothetical protein FLJ11729 /FL=gb:NM_024960.1	NM_024960		NP_705905
218827_s_at	0.016074	gb:NM_018069.1 /DEF=Homo sapiens hypothetical protein FLJ10352 (FLJ10352), mRNA. /FEA=mRNA /GEN=FLJ10352 /PROD=hypothetical protein FLJ10352 /DB_XREF=gi:8922371 /UG=Hs.100914 hypothetical protein FLJ10352 /FL=gb:NM_018069.1	NM_018069		

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
218829_s_at	1.26E-04	gb:NM_017780.1 /DEF=Homo sapiens hypothetical protein FLJ20357 (FLJ20357), mRNA. /FEA=mRNA /GEN=FLJ20357 /PROD=hypothetical protein FLJ20357 /DB_XREF=gi:8923329 /UG=Hs.105461 hypothetical protein FLJ20357. /FL=gb:NM_017780.1	NM_017780		
218833_at	3.09E-04	gb:NM_016653.1 /DEF=Homo sapiens sterile-alpha motif and leucine zipper containing kinase AZK (ZAK), mRNA. /FEA=mRNA /GEN=ZAK /PROD=sterile-alpha motif and leucine zippercontaining kinase AZK /DB_XREF=gi:7706600 /UG=Hs.115175 sterile-alpha motif and leucine zipper containing kinase AZK /FL=gb:AB049733.1 gb:AF238255.1 gb:AF251441.1 gb:NM_016653.1 gb:AB030034.1	NM_016653		NP_598407
218841_at	0.00161	gb:NM_024095.1 /DEF=Homo sapiens hypothetical protein MGC5540 (MGC5540), mRNA. /FEA=mRNA /GEN=MGC5540 /PROD=hypothetical protein MGC5540 /DB_XREF=gi:13129097 /UG=Hs.267400 hypothetical protein MGC5540 /FL=gb:BC001321.1 gb:NM_024095.1	NM_024095		NP_077000
218846_at	7.58E-04	gb:NM_004830.1 /DEF=Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 3 (130kD) (CRSP3), mRNA. /FEA=mRNA /GEN=CRSP3 /PROD=cofactor required for Sp1 transcriptionalactivation, subunit 3 (130kD) /DB_XREF=gi:7019352 /UG=Hs.29679 cofactor required for Sp1 transcriptional activation, subunit 3 (130kD) /FL=gb:AF135022.1 gb:AF105332.1 gb:NM_004830.1	NM_004830		NP_057063

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218850_s_at	0.020793	gb:NM_014240.1 /DEF=Homo sapiens LIM domains containing 1 (LIMD1), mRNA. /FEA=mRNA /GEN=LIMD1 /PROD=LIM domains containing 1 /DB_XREF=gi:7657306 /UG=Hs.48469 LIM domains containing 1 /FL=gb:NM_014240.1	NM_014240		NP_055055
218852_at	0.003171	gb:NM_017917.1 /DEF=Homo sapiens hypothetical protein FLJ20644 (FLJ20644), mRNA. /FEA=mRNA /GEN=FLJ20644 /PROD=hypothetical protein FLJ20644 /DB_XREF=gi:8923599 /UG=Hs.49376 hypothetical protein FLJ20644 /FL=gb:NM_017917.1	NM_017917		NP_060387
218853_s_at	0.004341	gb:NM_019556.1 /DEF=Homo sapiens hypothetical protein dJ473B4 (DJ473B4), mRNA. /FEA=mRNA /GEN=DJ473B4 /PROD=hypothetical protein dJ473B4 /DB_XREF=gi:9506542 /UG=Hs.57549 hypothetical protein dJ473B4 /FL=gb:NM_019556.1	NM_019556		NP_062456
218854_at	5.03E-04	gb:NM_013352.1 /DEF=Homo sapiens squamous cell carcinoma antigen recognized by T cell (SART-2), mRNA. /FEA=mRNA /GEN=SART-2 /PROD=squamous cell carcinoma antigen recognized by T cell /DB_XREF=gi:7019520 /UG=Hs.58636 squamous cell carcinoma antigen recognized by T cell /FL=gb:AF098066.1 gb:NM_013352.1	NM_013352		NP_037484

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218870_at	0.026891	gb:NM_018460.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM046 (BM046), mRNA. /FEA=mRNA /GEN=BM046 /PROD=uncharacterized bone marrow protein BM046 /DB_XREF=gi:8922105 /UG=Hs.177812 uncharacterized bone marrow protein BM046 /FL=gb:AF217507.1 gb:AF217522.1 gb:NM_018460.1 gb:AF212222.1	NM_018460		NP_060930
218871_x_at	2.05E-04	gb:NM_018590.1 /DEF=Homo sapiens hypothetical protein PRO0082 (PRO0082), mRNA. /FEA=mRNA /GEN=PRO0082 /PROD=hypothetical protein PRO0082 /DB_XREF=gi:8923965 /UG=Hs.180758 hypothetical protein PRO0082 /FL=gb:AF116646.1 gb:NM_018590.1	NM_018590		NP_061060
218872_at	2.75E-04	gb:NM_017899.1 /DEF=Homo sapiens hypothetical protein FLJ20607 (FLJ20607), mRNA. /FEA=mRNA /GEN=FLJ20607 /PROD=hypothetical protein FLJ20607 /DB_XREF=gi:8923562 /UG=Hs.18791 hypothetical protein FLJ20607 /FL=gb:NM_017899.1	NM_017899		NP_060369
218873_at	0.041055	gb:NM_017710.1 /DEF=Homo sapiens hypothetical protein FLJ20203 (FLJ20203), mRNA. /FEA=mRNA /GEN=FLJ20203 /PROD=hypothetical protein FLJ20203 /DB_XREF=gi:8923193 /UG=Hs.20594 hypothetical protein FLJ20203 /FL=gb:NM_017710.1	NM_017710		NP_115668
218880_at	0.010205	FOS-like antigen 2	N36408	Hs.301612	NP_005244

Gène Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218882_s_at	0.041055	gb:NM_006784.1 /DEF=Homo sapiens WD repeat domain 3 (WDR3), mRNA. /FEA=mRNA /GEN=WDR3 /PROD=WD repeat-containing protein 3 /DB_XREF=gi:5803220 /UG=Hs.33085 WD repeat domain 3 /FL=gb:AF083217.1 gb:NM_006784.1	NM_006784		NP_006775
218883_s_at	2.05E-04	gb:NM_024629.1 /DEF=Homo sapiens hypothetical protein FLJ23468 (FLJ23468), mRNA. /FEA=mRNA /GEN=FLJ23468 /PROD=hypothetical protein FLJ23468 /DB_XREF=gi:13375855 /UG=Hs.38178 hypothetical protein FLJ23468 /FL=gb:NM_024629.1	NM_024629		NP_078905
218888_s_at	0.048086	gb:NM_018092.1 /DEF=Homo sapiens hypothetical protein FLJ10430 (FLJ10430), mRNA. /FEA=mRNA /GEN=FLJ10430 /PROD=hypothetical protein FLJ10430 /DB_XREF=gi:8922419 /UG=Hs.6823 hypothetical protein FLJ10430 /FL=gb:NM_018092.1	NM_018092		NP_060562
218896_s_at	0.001116	gb:NM_018553.1 /DEF=Homo sapiens ELG protein (HSA277841), mRNA. /FEA=mRNA /GEN=HSA277841 /PROD=ELG protein /DB_XREF=gi:8923770 /UG=Hs.120963 ELG protein /FL=gb:NM_018553.1	NM_018553		NP_061023
218904_s_at	1.92E-04	gb:NM_017998.1 /DEF=Homo sapiens hypothetical protein FLJ10110 (FLJ10110), mRNA. /FEA=mRNA /GEN=FLJ10110 /PROD=hypothetical protein FLJ10110 /DB_XREF=gi:8922232 /UG=Hs.264363 hypothetical protein FLJ10110 /FL=gb:NM_017998.1	NM_017998		NP_060468

Gen Identifier	p-valu	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218909_at	0.027243	gb:NM_012424.1 /DEF=Homo sapiens ribosomal protein S6 kinase, 52kD, polypeptide 1 (RPS6KC1), mRNA. /FEA=mRNA /GEN=RPS6KC1 /PROD=ribosomal protein S6 kinase, 52kD, polypeptide1 /DB_XREF=gi:6912635 /UG=Hs.30352 ribosomal protein S6 kinase, 52kD, polypeptide 1 /FL=gb:AF037447.1 gb:NM_012424.1	NM_012424		NP_036556
218917_s_at	1.26E-04	gb:NM_018450.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM029 (BM029), mRNA. /FEA=mRNA /GEN=BM029 /PROD=uncharacterized bone marrow protein BM029 /DB_XREF=gi:8922086 /UG=Hs.8740 uncharacterized bone marrow protein BM029 /FL=gb:AF217506.1 gb:NM_018450.1	NM_018450		NP_624361
218921_at	0.022795	gb:NM_021805.1 /DEF=Homo sapiens single Ig IL-1R-related molecule (SIGIRR), mRNA. /FEA=mRNA /GEN=SIGIRR /PROD=single Ig IL-1R-related molecule /DB_XREF=gi:11141876 /UG=Hs.11809 single Ig IL-1R-related molecule /FL=gb:NM_021805.1 gb:BC003591.1	NM_021805		NP_068577
218923_at	0.001116	chitobiase, di-N-acetyl-	AW304174	Hs.135578	NP_004379
218930_s_at	0.00161	gb:NM_018374.1 /DEF=Homo sapiens hypothetical protein FLJ11273 (FLJ11273), mRNA. /FEA=mRNA /GEN=FLJ11273 /PROD=hypothetical protein FLJ11273 /DB_XREF=gi:8922965 /UG=Hs.3542 hypothetical protein FLJ11273 /FL=gb:NM_018374.1	NM_018374		NP_060844

Gen Id ntifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218937_at	0.033381	gb:NM_017810.1 /DEF=Homo sapiens hypothetical protein FLJ20417 (FLJ20417), mRNA. /FEA=mRNA /GEN=FLJ20417 /PROD=hypothetical protein FLJ20417 /DB_XREF=gi:8923385 /UG=Hs.10710 hypothetical protein FLJ20417 /FL=gb:NM_017810.1	NM_017810		NP_060280
218941_at	1.26E-04	gb:NM_012164.2 /DEF=Homo sapiens f-box and WD-40 domain protein 2 (FBXW2), mRNA. /FEA=mRNA /GEN=FBXW2 /PROD=f-box and WD-40 domain protein 2 /DB_XREF=gi:7549806 /UG=Hs.13755 f-box and WD-40 domain protein 2 /FL=gb:AF176698.1 gb:NM_012164.2	NM_012164		NP_036296
218942_at	0.002536	gb:NM_024779.1 /DEF=Homo sapiens hypothetical protein FLJ22055 (FLJ22055), mRNA. /FEA=mRNA /GEN=FLJ22055 /PROD=hypothetical protein FLJ22055 /DB_XREF=gi:13376135 /UG=Hs.144502 hypothetical protein FLJ22055 /FL=gb:NM_024779.1	NM_024779		NP_079055
218946_at	0.033381	gb:NM_015700.1 /DEF=Homo sapiens HIRIP5 protein (HIRIP5), mRNA. /FEA=mRNA /GEN=HIRIP5 /PROD=HIRIP5 protein /DB_XREF=gi:7661719 /UG=Hs.17368 HIRIP5 protein; HIRA-interacting protein 5. /FL=gb:AF132967.1 gb:NM_015700.1	NM_015700		NP_056515
218949_s_at	3.26E-04	gb:NM_018292.1 /DEF=Homo sapiens hypothetical protein FLJ10989 (FLJ10989), mRNA. /FEA=mRNA /GEN=FLJ10989 /PROD=hypothetical protein FLJ10989 /DB_XREF=gi:8922807 /UG=Hs.210778 hypothetical protein FLJ10989 /FL=gb:AL136679.1 gb:NM_018292.1	NM_018292		NP_060762

Gene Identifier	p-value	D scription	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218961_s_at	0.018693	gb:NM_007254.1 /DEF=Homo sapiens polynucleotide kinase 3-phosphatase (PNKP), mRNA. /FEA=mRNA /GEN=PNKP /PROD=polynucleotide kinase 3-phosphatase /DB_XREF=gi:6005835 /UG=Hs.78016 polynucleotide kinase 3-phosphatase /FL=gb:AF120499.1 gb:AF125807.1 gb:AF126486.1 gb:NM_007254.1	NM_007254		NP_009185
218963_s_at	0.021448	gb:NM_015515.1 /DEF=Homo sapiens DKFZP434G032 protein (DKFZP434G032), mRNA. /FEA=mRNA /GEN=DKFZP434G032 /PROD=DKFZP434G032 protein /DB_XREF=gi:7661573 /UG=Hs.9029 DKFZP434G032 protein /FL=gb:AF102848.1 gb:NM_015515.1	NM_015515		NP_775320
218964_at	0.021312	gb:NM_006465.1 /DEF=Homo sapiens dead ringer (Drosophila)-like 2 (bright and dead ringer) (DRIL2), mRNA. /FEA=mRNA /GEN=DRIL2 /PROD=dead ringer-like 2 /DB_XREF=gi:5453637 /UG=Hs.10431 dead ringer (Drosophila)-like 2 (bright and dead ringer) /FL=gb:AF116846.1 gb:NM_006465.1	NM_006465		NP_006456
218967_s_at	0.003597	gb:NM_030664.1 /DEF=Homo sapiens phosphotriesterase related (PTER), mRNA. /FEA=mRNA /GEN=PTER /PROD=phosphotriesterase related /DB_XREF=gi:13489055 /UG=Hs.129915 phosphotriesterase related /FL=gb:AF212237.1 gb:NM_030664.1	NM_030664		NP_109589

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218968_s_at	0.03018	gb:NM_018197.1 /DEF=Homo sapiens hypothetical protein FLJ10734 (FLJ10734), mRNA. /FEA=mRNA /GEN=FLJ10734 /PROD=hypothetical protein FLJ10734 /DB_XREF=gi:8922626 /UG=Hs.146332 hypothetical protein FLJ10734 /FL=gb:NM_018197.1	NM_018197		NP_071371
218969_at	0.023827	gb:NM_016069.1 /DEF=Homo sapiens CGI-136 protein (LOC51025), mRNA. /FEA=mRNA /GEN=LOC51025 /PROD=CGI-136 protein /DB_XREF=gi:7705633 /UG=Hs.16089 CGI-136 protein /FL=gb:BC005024.1 gb:AF151894.1 gb:NM_016069.1	NM_016069		NP_057153
218970_s_at	0.016934	gb:NM_015960.1 /DEF=Homo sapiens CGI-32 protein (LOC51076), mRNA. /FEA=mRNA /GEN=LOC51076 /PROD=CGI-32 protein /DB_XREF=gi:7705727 /UG=Hs.16606 CGI-32 protein /FL=gb:AF132966.1 gb:NM_015960.1	NM_015960		NP_057044
218979_at	0.001911	gb:NM_024945.1 /DEF=Homo sapiens hypothetical protein FLJ12888 (FLJ12888), mRNA. /FEA=mRNA /GEN=FLJ12888 /PROD=hypothetical protein FLJ12888 /DB_XREF=gi:13376426 /UG=Hs.284137 hypothetical protein FLJ12888 /FL=gb:NM_024945.1	NM_024945		NP_079221
218981_at	1.28E-04	gb:NM_020186.1 /DEF=Homo sapiens DC11 protein (DC11), mRNA. /FEA=mRNA /GEN=DC11 /PROD=DC11 protein /DB_XREF=gi:9910179 /UG=Hs.42785 DC11 protein /FL=gb:AF201933.1 gb:NM_020186.1	NM_020186		NP_064571

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218983_at	1.26E-04	gb:NM_016546.1 /DEF=Homo sapiens complement C1r-like proteinase precursor, (LOC51279), mRNA. /FEA=mRNA /GEN=LOC51279 /PROD=complement C1r-like proteinase precursor, /DB_XREF=gi:7706082 /UG=Hs.98571 complement C1r-like proteinase precursor, /FL=gb:AF178985.1 gb:NM_016546.1	NM_016546		NP_057630
218992_at	1.72E-04	gb:NM_018465.1 /DEF=Homo sapiens uncharacterized hematopoietic stemprogenitor cells protein MDS030 (MDS030), mRNA. /FEA=mRNA /GEN=MDS030 /PROD=uncharacterized hematopoietic stemprogenitorcells protein MDS030 /DB_XREF=gi:8923931 /UG=Hs.181385 uncharacterized hematopoietic stemprogenitor cells protein MDS030 /FL=gb:AF220050.1 gb:NM_018465.1 gb:AF225420.1	NM_018465		NP_060935
218996_at	0.001786	gb:NM_013342.1 /DEF=Homo sapiens TCF3 (E2A) fusion partner (in childhood Leukemia) (TFPT), mRNA. /FEA=mRNA /GEN=TFPT /PROD=TCF3 (E2A) fusion partner (in childhoodLeukemia) /DB_XREF=gi:7019370 /UG=Hs.233765 TCF3 (E2A) fusion partner (in childhood Leukemia) /FL=gb:BC001728.1 gb:BC004281.1 gb:AF052052.1 gb:NM_013342.1	NM_013342		NP_037474

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
218997_at	0.006705	gb:NM_022490.1 /DEF=Homo sapiens hypothetical protein FLJ13390 similar to PAF53 (FLJ13390), mRNA. /FEA=mRNA /GEN=FLJ13390 /PROD=hypothetical protein FLJ13390 similar to PAF53 /DB_XREF=gi:11968046 /UG=Hs.24884 hypothetical protein FLJ13390 similar to PAF53 /FL=gb:NM_022490.1 gb:BC001337.1	NM_022490		NP_071935
218998_at	1.26E-04	gb:NM_017832.1 /DEF=Homo sapiens hypothetical protein FLJ20457 (FLJ20457), mRNA. /FEA=mRNA /GEN=FLJ20457 /PROD=hypothetical protein FLJ20457 /DB_XREF=gi:8923430 /UG=Hs.29276 hypothetical protein FLJ20457 /FL=gb:AF153417.1 gb:NM_017832.1	NM_017832		NP_060302
219001_s_at	0.041055	gb:NM_024345.1 /DEF=Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA. /FEA=mRNA /GEN=MGC10765 /PROD=hypothetical protein MGC10765 /DB_XREF=gi:13236596 /UG=Hs.317589 hypothetical protein MGC10765 /FL=gb:BC003520.1 gb:NM_024345.1	NM_024345		NP_077321
219004_s_at	0.007581	gb:NM_018944.1 /DEF=Homo sapiens chromosome 21 open reading frame 45 (C21ORF45), mRNA. /FEA=mRNA /GEN=C21ORF45 /PROD=chromosome 21 open reading frame 45 /DB_XREF=gi:9506436 /UG=Hs.49932 chromosome 21 open reading frame 45 /FL=gb:NM_018944.1	NM_018944		NP_061817

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219012_s_at	0.006966	gb:NM_020193.1 /DEF=Homo sapiens GL002 protein (GL002), mRNA. /FEA=mRNA /GEN=GL002 /PROD=GL002 protein /DB_XREF=gi:9910245 /UG=Hs.9567 GL002 protein /FL=gb:AF226047.1 gb:NM_020193.1	NM_020193		NP_064578
219014_at	1.26E-04	gb:NM_016619.1 /DEF=Homo sapiens hypothetical protein (LOC51316), mRNA. /FEA=mRNA /GEN=LOC51316 /PROD=hypothetical protein /DB_XREF=gi:7706157 /UG=Hs.107139 hypothetical protein /FL=gb:AF208846.1 gb:NM_016619.1	NM_016619		NP_057703
219015_s_at	0.00228	gb:NM_018466.1 /DEF=Homo sapiens uncharacterized hematopoietic stemprogenitor cells protein MDS031 (MDS031), mRNA. /FEA=mRNA /GEN=MDS031 /PROD=uncharacterized hematopoietic stemprogenitorcells protein MDS031 /DB_XREF=gi:8923933 /UG=Hs.110853 uncharacterized hematopoietic stemprogenitor cells protein MDS031 /FL=gb:BC005336.1 gb:AF220051.1 gb:NM_018466.1	NM_018466		NP_060936
219016_at	1.36E-04	gb:NM_021826.1 /DEF=Homo sapiens hypothetical protein FLJ13149 (FLJ13149), mRNA. /FEA=mRNA /GEN=FLJ13149 /PROD=hypothetical protein FLJ13149 /DB_XREF=gi:11141902 /UG=Hs.112188 hypothetical protein FLJ13149 /FL=gb:NM_021826.1	NM_021826		NP_068598

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
219017_at	1.26E-04	gb:NM_018638.2 /DEF=Homo sapiens ethanolamine kinase (EKI1), mRNA. /FEA=mRNA /GEN=EKI1 /PROD=ethanolamine kinase /DB_XREF=gi:10092614 /UG=Hs.120439 ethanolamine kinase /FL=gb:NM_018638.2 gb:AF207600.2	NM_018638		NP_061108
219020_at	0.027418	gb:NM_022460.1 /DEF=Homo sapiens hypothetical protein FLJ14249 similar to HS1 binding protein 3 (FLJ14249), mRNA. /FEA=mRNA /GEN=FLJ14249 /PROD=hypothetical protein FLJ14249 similar to HS1 binding protein 3. /DB_XREF=gi:11968000 /UG=Hs.125520 hypothetical protein FLJ14249 similar to HS1 binding protein 3 /FL=gb:NM_022460.1	NM_022460		NP_567825
219035_s_at	1.26E-04	gb:NM_025126.1 /DEF=Homo sapiens hypothetical protein FLJ21786 (FLJ21786), mRNA. /FEA=mRNA /GEN=FLJ21786 /PROD=hypothetical protein FLJ21786 /DB_XREF=gi:13376704 /UG=Hs.316809 hypothetical protein FLJ21786 /FL=gb:NM_025126.1	NM_025126		NP_079402
219049_at	1.26E-04	gb:NM_018371.1 /DEF=Homo sapiens hypothetical protein FLJ11264 (FLJ11264), mRNA. /FEA=mRNA /GEN=FLJ11264 /PROD=hypothetical protein FLJ11264 /DB_XREF=gi:8922959 /UG=Hs.11260 hypothetical protein FLJ11264 /FL=gb:NM_018371.1	NM_018371		NP_060841

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
219065_s_at	0.010205	gb:NM_015955.1 /DEF=Homo sapiens CGI-27 protein (LOC51072), mRNA. /FEA=mRNA /GEN=LOC51072 /PROD=CGI-27 protein /DB_XREF=gi:7705719 /UG=Hs.20814 CGI-27 protein /FL=gb:AF132961.1 gb:NM_015955.1	NM_015955		NP_057039
219070_s_at	0.001483	gb:NM_023948.1 /DEF=Homo sapiens hypothetical protein AF053356_CDS3 (AF053356_CDS3), mRNA. /FEA=mRNA /GEN=AF053356_CDS3 /PROD=AF053356_CDS3 protein /DB_XREF=gi:13122607 /UG=Hs.296200 hypothetical protein AF053356_CDS3 /FL=gb:NM_023948.1 gb:BC005042.1	NM_023948		NP_076438
219074_at	0.036688	gb:NM_018241.1 /DEF=Homo sapiens hypothetical protein FLJ10846 (FLJ10846), mRNA. /FEA=mRNA /GEN=FLJ10846 /PROD=hypothetical protein FLJ10846 /DB_XREF=gi:8922706 /UG=Hs.32271 hypothetical protein FLJ10846 /FL=gb:NM_018241.1	NM_018241		NP_060711
219079_at	1.26E-04	gb:NM_016230.1 /DEF=Homo sapiens flavohemoprotein b5+b5R (LOC51167), mRNA. /FEA=mRNA /GEN=LOC51167 /PROD=flavohemoprotein b5+b5R /DB_XREF=gi:7705898 /UG=Hs.5741 flavohemoprotein b5+b5R /FL=gb:AF169803.1 gb:NM_016230.1	NM_016230		NP_057314
219099_at	0.010205	gb:NM_020375.1 /DEF=Homo sapiens chromosome 12 open reading frame 5 (C12ORF5), mRNA. /FEA=mRNA /GEN=C12ORF5 /PROD=chromosome 12 open reading frame 5 /DB_XREF=gi:9966848 /UG=Hs.24792 chromosome 12 open reading frame 5 /FL=gb:NM_020375.1	NM_020375		NP_065108

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219104_at	5.03E-04	gb:NM_016422.1 /DEF=Homo sapiens C3HC4-like zinc finger protein (ZFP26), mRNA. /FEA=mRNA /GEN=ZFP26 /PROD=C3HC4-like zinc finger protein /DB_XREF=gi:7706776 /UG=Hs.44685 C3HC4-like zinc finger protein /FL=gb:AF214680.1 gb:NM_016422.1	NM_016422		NP_057506
219112_at	0.00161	gb:NM_016340.1 /DEF=Homo sapiens PDZ domain-containing guanine nucleotide exchange factor 1 (LOC51735), mRNA. /FEA=mRNA /GEN=LOC51735 /PROD=PDZ domain-containing guanine nucleotide exchange factor 1 /DB_XREF=gi:7706512 /UG=Hs.174795 PDZ domain-containing guanine nucleotide exchange factor 1 /FL=gb:AF117947.1 gb:NM_016340.1	NM_016340		NP_057424
219116_s_at	0.003395	gb:NM_018185.1 /DEF=Homo sapiens hypothetical protein FLJ10704 (FLJ10704), mRNA. /FEA=mRNA /GEN=FLJ10704 /PROD=hypothetical protein FLJ10704 /DB_XREF=gi:8922602 /UG=Hs.23618 hypothetical protein FLJ10704 /FL=gb:NM_018185.1	NM_018185		NP_060655
219123_at	0.022926	gb:NM_014519.1 /DEF=Homo sapiens zinc finger protein 232 (ZNF232), mRNA. /FEA=mRNA /GEN=ZNF232 /PROD=zinc finger protein 232 /DB_XREF=gi:7657704 /UG=Hs.279914 zinc finger protein 232 /FL=gb:NM_014519.1	NM_014519		NP_055334

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219129_s_at	1.49E-04	gb:NM_024632.1 /DEF=Homo sapiens hypothetical protein FLJ11526 (FLJ11526), mRNA. /FEA=mRNA /GEN=FLJ11526 /PROD=hypothetical protein FLJ11526 /DB_XREF=gi:13375861 /UG=Hs.38750 hypothetical protein FLJ11526 /FL=gb:NM_024632.1	NM_024632		NP_078908
219132_at	5.03E-04	gb:NM_021255.1 /DEF=Homo sapiens pellino (Drosophila) homolog 2 (PELI2), mRNA. /FEA=mRNA /GEN=PELI2 /PROD=pellino (Drosophila) homolog 2 /DB_XREF=gi:10864062 /UG=Hs.44038 pellino (Drosophila) homolog 2 /FL=gb:AF302502.1 gb:NM_021255.1	NM_021255		NP_067078
219137_s_at	1.26E-04	gb:NM_020194.1 /DEF=Homo sapiens GL004 protein (GL004), mRNA. /FEA=mRNA /GEN=GL004 /PROD=GL004 protein /DB_XREF=gi:9910247 /UG=Hs.7045 GL004 protein /FL=gb:AF226049.1 gb:NM_020194.1	NM_020194		NP_064579
219149_x_at	0.004908	gb:NM_016216.1 /DEF=Homo sapiens RNA lariat debranching enzyme (LOC51163), mRNA. /FEA=mRNA /GEN=LOC51163 /PROD=RNA lariat debranching enzyme /DB_XREF=gi:7705890 /UG=Hs.118194 RNA lariat debranching enzyme /FL=gb:AF180919.2 gb:NM_016216.1	NM_016216		NP_057300
219156_at	1.26E-04	gb:NM_018373.1 /DEF=Homo sapiens hypothetical protein FLJ11271 (FLJ11271), mRNA. /FEA=mRNA /GEN=FLJ11271 /PROD=hypothetical protein FLJ11271 /DB_XREF=gi:8922963 /UG=Hs.109654 hypothetical protein FLJ11271 /FL=gb:NM_018373.1	NM_018373		NP_060843

Gen Identifier	p-value	D scription	Gene Accession No.	Unigen Accession No.	Prot in Acc ssion No.
219157_at	0.001116	gb:NM_007246.1 /DEF=Homo sapiens kelch (Drosophila)-like 2 (Mayven) (KLHL2), mRNA. /FEA=mRNA /GEN=KLHL2 /PROD=kelch (Drosophila)-like 2 (Mayven) /DB_XREF=gi:6005702 /UG=Hs.122967 kelch (Drosophila)-like 2 (Mayven) /FL=gb:AF059569.1 gb:NM_007246.1	NM_007246		NP_009177
219158_s_at	6.68E-04	gb:NM_025085.1 /DEF=Homo sapiens hypothetical protein FLJ13340 (FLJ13340), mRNA. /FEA=mRNA /GEN=FLJ13340 /PROD=hypothetical protein FLJ13340 /DB_XREF=gi:13376639 /UG=Hs.125034 hypothetical protein FLJ13340 /FL=gb:AF327722.1 gb:NM_025085.1	NM_025085		NP_476516
219161_s_at	0.004341	gb:NM_016951.2 /DEF=Homo sapiens transmembrane proteolipid (HSPC224), mRNA. /FEA=mRNA /GEN=HSPC224 /PROD=transmembrane proteolipid /DB_XREF=gi:10092593 /UG=Hs.15159 chemokine-like factor, alternatively spliced /FL=gb:NM_016951.2 gb:BC004380.1 gb:AF057306.1 gb:AF151058.1 gb:AF135380.2	NM_016951		NP_058647
219163_at	0.007052	gb:NM_017656.1 /DEF=Homo sapiens hypothetical protein FLJ20079 (FLJ20079), mRNA. /FEA=mRNA /GEN=FLJ20079 /PROD=hypothetical protein FLJ20079 /DB_XREF=gi:8923085 /UG=Hs.165948 hypothetical protein FLJ20079 /FL=gb:NM_017656.1	NM_017656		NP_060126

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219165_at	0.016934	gb:NM_021630.1 /DEF=Homo sapiens PDZ-LIM protein mystique (LOC59346), mRNA. /FEA=mRNA /GEN=LOC59346 /PROD=PDZ-LIM protein mystique /DB_XREF=gi:11055999 /UG=Hs.19447 PDZ-LIM protein mystique /FL=gb:AY007729.1 gb:NM_021630.1	NM_021630		NP_789847
219172_at	0.034221	gb:NM_024954.1 /DEF=Homo sapiens hypothetical protein FLJ11807 (FLJ11807), mRNA. /FEA=mRNA /GEN=FLJ11807 /PROD=hypothetical protein FLJ11807 /DB_XREF=gi:13376438 /UG=Hs.285813 hypothetical protein FLJ11807 /FL=gb:NM_024954.1	NM_024954		NP_079230
219180_s_at	0.02145	hypothetical protein FLJ20695	AI817074	Hs.5420	NP_060399
219190_s_at	0.005852	gb:NM_017629.1 /DEF=Homo sapiens hypothetical protein FLJ20033 (FLJ20033), mRNA. /FEA=mRNA /GEN=FLJ20033 /PROD=hypothetical protein FLJ20033 /DB_XREF=gi:8923033 /UG=Hs.134757 hypothetical protein FLJ20033 /FL=gb:NM_017629.1	NM_017629		NP_060099
219191_s_at	3.26E-04	gb:NM_016293.1 /DEF=Homo sapiens bridging integrator 2 (BIN2), mRNA. /FEA=mRNA /GEN=BIN2 /PROD=breast cancer associated protein BRAP1 /DB_XREF=gi:7706486 /UG=Hs.14770 bridging integrator 2 /FL=gb:AF146531.1 gb:AB032698.1 gb:NM_016187.1 gb:NM_016293.1	NM_016293		NP_057377
219221_at	0.003539	gb:NM_024724.1 /DEF=Homo sapiens hypothetical protein FLJ22332 (FLJ22332), mRNA. /FEA=mRNA /GEN=FLJ22332 /PROD=hypothetical protein FLJ22332 /DB_XREF=gi:13376033 /UG=Hs.111092 hypothetical protein FLJ22332 /FL=gb:NM_024724.1	NM_024724		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219228_at	6.68E-04	gb:NM_018555.2 /DEF=Homo sapiens C2H2-like zinc finger protein (ZNF361), mRNA. /FEA=mRNA /GEN=ZNF361 /PROD=C2H2-like zinc finger protein /DB_XREF=gi:10092612 /UG=Hs.147644 zinc finger protein 331 /FL=gb:AF251515.2 gb:NM_018555.2 gb:AF272148.1	NM_018555		NP_061025
219229_at	0.001116	gb:NM_013272.2 /DEF=Homo sapiens solute carrier family 21 (organic anion transporter), member 11 (SLC21A11), mRNA. /FEA=mRNA /GEN=SLC21A11 /PROD=solute carrier family 21 (organic anion transporter), member 11 /DB_XREF=gi:7706713 /UG=Hs.14805 solute carrier family 21 (organic anion transporter), member 11 /FL=gb:AF205074.1 gb:AF187816.1 gb:AB031050.2 gb:NM_013272.2	NM_013272		NP_037404
219232_s_at	0.014208	gb:NM_022073.1 /DEF=Homo sapiens hypothetical protein FLJ21620 (FLJ21620), mRNA. /FEA=mRNA /GEN=FLJ21620 /PROD=hypothetical protein FLJ21620 /DB_XREF=gi:11545786 /UG=Hs.18878 hypothetical protein FLJ21620 /FL=gb:NM_022073.1	NM_022073		NP_203130
219238_at	0.006966	gb:NM_017837.1 /DEF=Homo sapiens hypothetical protein FLJ20477 (FLJ20477), mRNA. /FEA=mRNA /GEN=FLJ20477 /PROD=hypothetical protein FLJ20477 /DB_XREF=gi:8923441 /UG=Hs.26994 hypothetical protein FLJ20477 /FL=gb:NM_017837.1	NM_017837		NP_060307

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219253_at	1.26E-04	gb:NM_024121.1 /DEF=Homo sapiens hypothetical protein FLJ20979 (FLJ20979), mRNA. /FEA=mRNA /GEN=FLJ20979 /PROD=hypothetical protein FLJ20979 /DB_XREF=gi:13129145 /UG=Hs.44680 hypothetical protein FLJ20979 /FL=gb:NM_024121.1	NM_024121		
219259_at	0.00228	gb:NM_022367.1 /DEF=Homo sapiens hypothetical protein FLJ12287 similar to semaphorins (FLJ12287), mRNA. /FEA=mRNA /GEN=FLJ12287 /PROD=hypothetical protein FLJ12287 similar to semaphorins /DB_XREF=gi:11641290 /UG=Hs.7634 hypothetical protein FLJ12287 similar to semaphorins /FL=gb:NM_022367.1 gb:AB029394.1	NM_022367		NP_071762
219266_at	0.02145	gb:NM_021632.1 /DEF=Homo sapiens zinc-finger protein ZBRK1 (ZBRK1), mRNA. /FEA=mRNA /GEN=ZBRK1 /PROD=zinc-finger protein ZBRK1 /DB_XREF=gi:11056003 /UG=Hs.130965 zinc-finger protein ZBRK1 /FL=gb:AF295096.1 gb:AF309561.1 gb:NM_021632.1	NM_021632		NP_067645
219281_at	1.26E-04	gb:NM_012331.2 /DEF=Homo sapiens methionine sulfoxide reductase A (MSRA), mRNA. /FEA=mRNA /GEN=MSRA /PROD=methionine sulfoxide reductase A /DB_XREF=gi:13259538 /UG=Hs.26458 methionine sulfoxide reductase A /FL=gb:NM_012331.2 gb:AF183420.1	NM_012331		NP_036463

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219283_at	1.26E-04	gb:NM_014158.1 /DEF=Homo sapiens HSPC067 protein (HSPC067), mRNA. /FEA=mRNA /GEN=HSPC067 /PROD=HSPC067 protein /DB_XREF=gi:7661771 /UG=Hs.279938 HSPC067 protein /FL=gb:AF161552.1 gb:NM_014158.1	NM_014158		
219292_at	3.56E-04	gb:NM_018105.1 /DEF=Homo sapiens hypothetical protein FLJ10477 (FLJ10477), mRNA. /FEA=mRNA /GEN=FLJ10477 /PROD=hypothetical protein FLJ10477 /DB_XREF=gi:8922445 /UG=Hs.7432 hypothetical protein FLJ10477 /FL=gb:NM_018105.1	NM_018105		NP_060575
219297_at	0.026891	gb:NM_019045.1 /DEF=Homo sapiens similar to rab11-binding protein (FLJ11116), mRNA. /FEA=mRNA /GEN=FLJ11116 /PROD=similar to rab11-binding protein /DB_XREF=gi:9506636 /UG=Hs.98510 similar to rab11-binding protein /FL=gb:NM_019045.1	NM_019045		NP_061918
219304_s_at	0.041055	gb:NM_025208.1 /DEF=Homo sapiens spinal cord-derived growth factor-B (SCDGF-B), mRNA. /FEA=mRNA /GEN=SCDGF-B /PROD=spinal cord-derived growth factor-B /DB_XREF=gi:13376807 /UG=Hs.112885 spinal cord-derived growth factor-B /FL=gb:AB033832.1 gb:AF113216.1 gb:NM_025208.1 gb:AY027517.1	NM_025208		NP_149126
219322_s_at	0.02145	gb:NM_017818.1 /DEF=Homo sapiens WD repeat domain 8 (WDR8), mRNA. /FEA=mRNA /GEN=WDR8 /PROD=WD repeat domain 8 /DB_XREF=gi:8923402 /UG=Hs.31714 hypothetical protein FLJ20430 /FL=gb:AB034912.1 gb:NM_017818.1	NM_017818		NP_060288

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
219324_at	0.001173	gb:NM_024313.1 /DEF=Homo sapiens hypothetical protein MGC3731 (MGC3731), mRNA. /FEA=mRNA /GEN=MGC3731 /PROD=hypothetical protein MGC3731 /DB_XREF=gi:13236552 /UG=Hs.326372 hypothetical protein MGC3731 /FL=gb:BC002808.1 gb:NM_024313.1	NM_024313		NP_077289
219326_s_at	3.93E-04	gb:NM_006577.2 /DEF=Homo sapiens beta-1,3-N-acetylglucosaminyltransferase (BETA3GNT), mRNA. /FEA=mRNA /GEN=BETA3GNT /PROD=beta-1,3-N-acetylglucosaminyltransferase /DB_XREF=gi:9845237 /UG=Hs.48730 beta-1,3-N-acetylglucosaminyltransferase bGnT-2 /FL=gb:AF288208.1 gb:AF092051.2 gb:NM_006577.2	NM_006577		NP_150274
219329_s_at	0.007779	gb:NM_016085.1 /DEF=Homo sapiens apoptosis related protein APR-3 (APR-3), mRNA. /FEA=mRNA /GEN=APR-3 /PROD=apoptosis related protein APR-3 /DB_XREF=gi:7706360 /UG=Hs.9527 apoptosis related protein APR-3 /FL=gb:AF144055.2 gb:NM_016085.1	NM_016085		NP_542159
219336_s_at	0.034317	gb:NM_015947.1 /DEF=Homo sapiens CGI-18 protein (LOC51008), mRNA. /FEA=mRNA /GEN=LOC51008 /PROD=CGI-18 protein /DB_XREF=gi:7705601 /UG=Hs.121599 CGI-18 protein /FL=gb:AY013290.1 gb:AF132952.1 gb:NM_015947.1	NM_015947		NP_057031

Gene Identifier	p-value	D scription	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219356_s_at	0.001116	gb:NM_016410.1 /DEF=Homo sapiens hypothetical protein (HSPC177), mRNA. /FEA=mRNA /GEN=HSPC177 /PROD=hypothetical protein /DB_XREF=gi:7705488 /UG=Hs.279777 hypothetical protein /FL=gb:AF161525.1 gb:NM_016410.1	NM_016410		NP_057494
219357_at	0.007558	gb:NM_014027.1 /DEF=Homo sapiens HSPC018 protein (HSPC018), mRNA. /FEA=mRNA /GEN=HSPC018 /PROD=HSPC018 protein /DB_XREF=gi:7661735 /UG=Hs.279853 HSPC018 protein /FL=gb:BC001701.1 gb:AF077204.1 gb:NM_014027.1	NM_014027		NP_054746
219359_at	3.26E-04	gb:NM_025092.1 /DEF=Homo sapiens hypothetical protein FLJ22635 (FLJ22635), mRNA. /FEA=mRNA /GEN=FLJ22635 /PROD=hypothetical protein FLJ22635 /DB_XREF=gi:13376651 /UG=Hs.288529 hypothetical protein FLJ22635 /FL=gb:NM_025092.1	NM_025092		NP_079368
219371_s_at	0.005852	gb:NM_016270.1 /DEF=Homo sapiens Kruppel-like factor (LOC51713), mRNA. /FEA=mRNA /GEN=LOC51713 /PROD=Kruppel-like factor /DB_XREF=gi:7706468 /UG=Hs.107740 Kruppel-like factor 2 (lung) /FL=gb:AF134053.1 gb:AF205849.1 gb:NM_016270.1	NM_016270		NP_057354
219384_s_at	1.72E-04	gb:NM_012091.2 /DEF=Homo sapiens adenosine deaminase, tRNA-specific 1 (ADAT1), mRNA. /FEA=mRNA /GEN=ADAT1 /PROD=adenosine deaminase, tRNA-specific 1 /DB_XREF=gi:7382475 /UG=Hs.188661 adenosine deaminase, tRNA-specific 1 /FL=gb:BC002758.1 gb:AF125188.1 gb:NM_012091.2	NM_012091		NP_036223

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219394_at	0.00161	gb:NM_024419.1 /DEF=Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA. /FEA=mRNA /GEN=PGS1 /PROD=Phosphatidylglycerophosphate Synthase /DB_XREF=gi:13259369 /UG=Hs.278682 Phosphatidylglycerophosphate Synthase /FL=gb:NM_024419.1	NM_024419		NP_077733
219396_s_at	0.021312	gb:NM_024608.1 /DEF=Homo sapiens hypothetical protein FLJ22402 (FLJ22402), mRNA. /FEA=mRNA /GEN=FLJ22402 /PROD=hypothetical protein FLJ22402 /DB_XREF=gi:13375816 /UG=Hs.28355 hypothetical protein FLJ22402 /FL=gb:NM_024608.1	NM_024608		NP_078884
219397_at	0.003171	gb:NM_025147.1 /DEF=Homo sapiens hypothetical protein FLJ13448 (FLJ13448), mRNA. /FEA=mRNA /GEN=FLJ13448 /PROD=hypothetical protein FLJ13448 /DB_XREF=gi:13376736 /UG=Hs.288945 hypothetical protein FLJ13448 /FL=gb:NM_025147.1	NM_025147		NP_079423
219403_s_at	1.41E-04	gb:NM_006665.1 /DEF=Homo sapiens heparanase (HPSE), mRNA. /FEA=mRNA /GEN=HPSE /PROD=heparanase /DB_XREF=gi:5729872 /UG=Hs.44227 heparanase /FL=gb:AF165154.1 gb:AF152376.1 gb:NM_006665.1 gb:AF084467.1 gb:AF155510.1	NM_006665		NP_006656
219405_at	0.023856	gb:NM_018073.1 /DEF=Homo sapiens hypothetical protein FLJ10369 (FLJ10369), mRNA. /FEA=mRNA /GEN=FLJ10369 /PROD=hypothetical protein FLJ10369 /DB_XREF=gi:8922379 /UG=Hs.53940 hypothetical protein FLJ10369 /FL=gb:NM_018073.1	NM_018073		NP_060543

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219409_at	0.046189	gb:NM_024700.1 /DEF=Homo sapiens hypothetical protein FLJ12553 (FLJ12553), mRNA. /FEA=mRNA /GEN=FLJ12553 /PROD=hypothetical protein FLJ12553 /DB_XREF=gi:13375986 /UG=Hs.99601 hypothetical protein FLJ12553 /FL=gb:NM_024700.1	NM_024700		NP_078976
219410_at	0.037692	gb:NM_018004.1 /DEF=Homo sapiens hypothetical protein FLJ10134 (FLJ10134), mRNA. /FEA=mRNA /GEN=FLJ10134 /PROD=hypothetical protein FLJ10134 /DB_XREF=gi:8922242 /UG=Hs.104800 hypothetical protein FLJ10134 /FL=gb:NM_018004.1	NM_018004		NP_060474
219426_at	7.58E-04	gb:NM_024852.1 /DEF=Homo sapiens hypothetical protein FLJ12765 (FLJ12765), mRNA. /FEA=mRNA /GEN=FLJ12765 /PROD=hypothetical protein FLJ12765 /DB_XREF=gi:13376274 /UG=Hs.189829 hypothetical protein FLJ12765 /FL=gb:NM_024852.1	NM_024852		NP_803171
219444_at	1.26E-04	gb:NM_021946.1 /DEF=Homo sapiens hypothetical protein FLJ11362 (FLJ11362), mRNA. /FEA=mRNA /GEN=FLJ11362 /PROD=hypothetical protein FLJ11362 /DB_XREF=gi:11345489 /UG=Hs.8929 hypothetical protein FLJ11362 /FL=gb:NM_021946.1	NM_021946		NP_068765

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219445_at	1.49E-04	gb:NM_015711.1 /DEF=Homo sapiens glioma tumor suppressor candidate region gene 1 (GLTSCR1), mRNA. /FEA=mRNA /GEN=GLTSCR1 /PROD=glioma tumor suppressor candidate region gene 1 /DB_XREF=gi:7657127 /UG=Hs.97244 glioma tumor suppressor candidate region gene 1 /FL=gb:AF182077.1 gb:NM_015711.1	NM_015711		NP_056526
219446_at	0.042466	gb:NM_018157.1 /DEF=Homo sapiens hypothetical protein FLJ10620 (FLJ10620), mRNA. /FEA=mRNA /GEN=FLJ10620 /PROD=hypothetical protein FLJ10620 /DB_XREF=gi:8922554 /UG=Hs.99445 hypothetical protein FLJ10620 /FL=gb:NM_018157.1	NM_018157		NP_060627
219449_s_at	3.26E-04	gb:NM_017866.1 /DEF=Homo sapiens hypothetical protein FLJ20533 (FLJ20533), mRNA. /FEA=mRNA /GEN=FLJ20533 /PROD=hypothetical protein FLJ20533 /DB_XREF=gi:8923499 /UG=Hs.106650 hypothetical protein FLJ20533 /FL=gb:BC002748.1 gb:NM_017866.1	NM_017866		NP_060336
219457_s_at	0.005039	gb:NM_024832.1 /DEF=Homo sapiens hypothetical protein FLJ22439 (FLJ22439), mRNA. /FEA=mRNA /GEN=FLJ22439 /PROD=hypothetical protein FLJ22439 /DB_XREF=gi:13376237 /UG=Hs.180040 hypothetical protein FLJ22439 /FL=gb:NM_024832.1	NM_024832		NP_079108

Gen Identifier	p-value	Description	Gene Accession No.	Unig ne Accession No.	Protein Accession No.
219458_s_at	0.026891	gb:NM_022072.1 /DEF=Homo sapiens hypothetical protein FLJ22609 (FLJ22609), mRNA. /FEA=mRNA /GEN=FLJ22609 /PROD=hypothetical protein FLJ22609 /DB_XREF=gi:11545784 /UG=Hs.18740 hypothetical protein FLJ22609 /FL=gb:NM_022072.1	NM_022072		NP_071355
219459_at	0.03002	gb:NM_018082.1 /DEF=Homo sapiens hypothetical protein FLJ10388 (FLJ10388), mRNA. /FEA=mRNA /GEN=FLJ10388 /PROD=hypothetical protein FLJ10388 /DB_XREF=gi:8922398 /UG=Hs.197642 hypothetical protein FLJ10388 /FL=gb:NM_018082.1	NM_018082		NP_060552
219460_s_at	1.28E-04	gb:NM_017849.1 /DEF=Homo sapiens hypothetical protein FLJ20507 (FLJ20507), mRNA. /FEA=mRNA /GEN=FLJ20507 /PROD=hypothetical protein FLJ20507 /DB_XREF=gi:8923465 /UG=Hs.202955 hypothetical protein FLJ20507 /FL=gb:NM_017849.1	NM_017849		NP_060319
219467_at	0.033381	gb:NM_017676.1 /DEF=Homo sapiens hypothetical protein FLJ20125 (FLJ20125), mRNA. /FEA=mRNA /GEN=FLJ20125 /PROD=hypothetical protein FLJ20125 /DB_XREF=gi:8923123 /UG=Hs.24088 hypothetical protein FLJ20125 /FL=gb:NM_017676.1	NM_017676		NP_060146

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219485_s_at	0.001116	gb:NM_002814.1 /DEF=Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 10 (PSMD10), mRNA. /FEA=mRNA /GEN=PSMD10 /PROD=proteasome (prosome, macropain) 26S subunit, non-ATPase, 10 /DB_XREF=gi:4506216 /UG=Hs.7756 proteasome (prosome, macropain) 26S subunit, non-ATPase, 10 /FL=gb:AB009619.1 gb:D83197.1 gb:NM_002814.1	NM_002814		NP_736606
219492_at	3.26E-04	gb:NM_012110.1 /DEF=Homo sapiens cystein-rich hydrophobic domain 2 (CHIC2), mRNA. /FEA=mRNA /GEN=CHIC2 /PROD=cystein-rich hydrophobic domain 2 /DB_XREF=gi:6912275 /UG=Hs.119488 cystein-rich hydrophobic domain 2 /FL=gb:AF159423.1 gb:NM_012110.1	NM_012110		NP_036242
219493_at	0.005379	gb:NM_024745.1 /DEF=Homo sapiens hypothetical protein FLJ22009 (FLJ22009), mRNA. /FEA=mRNA /GEN=FLJ22009 /PROD=hypothetical protein FLJ22009 /DB_XREF=gi:13376069 /UG=Hs.123253 hypothetical protein FLJ22009 /FL=gb:NM_024745.1	NM_024745		NP_079021
219495_s_at	0.046329	gb:NM_013256.1 /DEF=Homo sapiens zinc finger protein 180 (HHZ168) (ZNF180), mRNA. /FEA=mRNA /GEN=ZNF180 /PROD=zinc finger protein 180 (HHZ168) /DB_XREF=gi:7019578 /UG=Hs.130683 zinc finger protein 180 (HHZ168) /FL=gb:AF192913.1 gb:NM_013256.1	NM_013256		NP_037388

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219497_s_at	0.013088	gb:NM_022893.1 /DEF=Homo sapiens B-cell CLLlymphoma 11A (zinc finger protein) (BCL11A); mRNA. /FEA=mRNA /GEN=BCL11A /PROD=B-cell CLLlymphoma 11A (zinc finger protein) /DB_XREF=gi:13375629 /UG=Hs.130881 B-cell CLLlymphoma 11A (zinc finger protein) /FL=gb:NM_022893.1 gb:NM_018014.1	NM_022893		NP_612569
219502_at	0.046189	gb:NM_018248.1 /DEF=Homo sapiens hypothetical protein FLJ10858 (FLJ10858), mRNA. /FEA=mRNA /GEN=FLJ10858 /PROD=hypothetical protein FLJ10858 /DB_XREF=gi:8922721 /UG=Hs.134403 hypothetical protein FLJ10858 /FL=gb:NM_018248.1	NM_018248		NP_060718
219515_at	0.013223	gb:NM_020228.1 /DEF=Homo sapiens PR domain containing 10 (PRDM10), mRNA. /FEA=mRNA /GEN=PRDM10 /PROD=PR domain containing 10 /DB_XREF=gi:9910503 /UG=Hs.275086 PR domain containing 10 /FL=gb:AF275817.1 gb:NM_020228.1	NM_020228		NP_064613
219520_s_at	0.033381	gb:NM_018458.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM042 (BM042), mRNA. /FEA=mRNA /GEN=BM042 /PROD=uncharacterized bone marrow protein BM042 /DB_XREF=gi:8922101 /UG=Hs.324136 uncharacterized bone marrow protein BM042 /FL=gb:AF217518.1 gb:NM_018458.1	NM_018458		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
219528_s_at	0.005852	gb:NM_022898.1 /DEF=Homo sapiens B-cell lymphomaleukaemia 11B (BCL11B), mRNA. /FEA=mRNA /GEN=BCL11B /PROD=B-cell lymphomaleukaemia 11B /DB_XREF=gi:12597634 /UG=Hs.57987 B-cell CLLlymphoma 11B (zinc finger protein) /FL=gb:NM_022898.1	NM_022898		NP_612808
219544_at	0.00228	gb:NM_024808.1 /DEF=Homo sapiens hypothetical protein FLJ22624 (FLJ22624), mRNA. /FEA=mRNA /GEN=FLJ22624 /PROD=hypothetical protein FLJ22624 /DB_XREF=gi:13376190 /UG=Hs.166425 hypothetical protein FLJ22624. /FL=gb:NM_024808.1	NM_024808		NP_079084
219547_at	1.26E-04	gb:NM_004376.1 /DEF=Homo sapiens COX15 (yeast) homolog, cytochrome c oxidase assembly protein (COX15), mRNA. /FEA=mRNA /GEN=COX15 /PROD=COX15 (yeast) homolog, cytochrome c oxidase assembly protein /DB_XREF=gi:4758035 /UG=Hs.226581 COX15 (yeast) homolog, cytochrome c oxidase assembly protein /FL=gb:AF044323.1 gb:NM_004376.1	NM_004376		NP_510870
219549_s_at	7.58E-04	gb:NM_006054.1 /DEF=Homo sapiens reticulon 3 (RTN3), mRNA. /FEA=mRNA /GEN=RTN3 /PROD=reticulon 3 /DB_XREF=gi:5174654 /UG=Hs.252831 reticulon 3 /FL=gb:BC000634.1 gb:AF059524.1 gb:AF119297.1 gb:NM_006054.1	NM_006054		NP_006045

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219551_at	1.32E-04	gb:NM_018456.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM040 (BM040), mRNA. /FEA=mRNA /GEN=BM040 /PROD=uncharacterized bone marrow protein BM040 /DB_XREF=gi:8922098 /UG=Hs.26892 uncharacterized bone marrow protein BM040 /FL=gb:AF217516.1 gb:NM_018456.1	NM_018456		NP_060926
219563_at	0.004341	gb:NM_024633.1 /DEF=Homo sapiens hypothetical protein FLJ21276 (FLJ21276), mRNA. /FEA=mRNA /GEN=FLJ21276 /PROD=hypothetical protein FLJ21276 /DB_XREF=gi:13375863 /UG=Hs.41502 hypothetical protein FLJ21276 /FL=gb:NM_024633.1	NM_024633		NP_078909
219571_s_at	2.05E-04	gb:NM_016265.1 /DEF=Homo sapiens GIOT-3 for gonadotropin inducible transcription repressor-3 (GIOT-3), mRNA. /FEA=mRNA /GEN=GIOT-3 /PROD=GIOT-3 for gonadotropin inducible transcription repressor-3 /DB_XREF=gi:7706464 /UG=Hs.102397 GIOT-3 for gonadotropin inducible transcription repressor-3 /FL=gb:AB021643.1 gb:NM_016265.1	NM_016265		NP_057349
219575_s_at	0.010205	gb:NM_022341.1 /DEF=Homo sapiens peptide deformylase-like protein (LOC64146), mRNA. /FEA=mRNA /GEN=LOC64146 /PROD=peptide deformylase-like protein /DB_XREF=gi:11641242 /UG=Hs.130849 peptide deformylase-like protein /FL=gb:AF239156.1 gb:NM_022341.1 gb:AF322879.1	NM_022341		NP_071736

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219594_at	1.72E-04	gb:NM_016533.1 /DEF=Homo sapiens ninjurin 2 (NINJ2), mRNA /FEA=mRNA /GEN=NINJ2 /PROD=ninjurin 2 /DB_XREF=gi:7706622 /UG=Hs.239208. ninjurin 2 /FL=gb:AF205633.1 gb:NM_016533.1	NM_016533		
219607_s_at	3.56E-04	gb:NM_024021.1 /DEF=Homo sapiens membrane-spanning 4-domains, subfamily A, member 4 (MS4A4), mRNA /FEA=mRNA /GEN=MS4A4 /PROD=membrane-spanning 4-domains, subfamily A, member 4 /DB_XREF=gi:13430865 /UG=Hs.325960 membrane-spanning 4-domains, subfamily A, member 4 /FL=gb:AB013102.1 gb:NM_024021.1 gb:AF068288.1 gb:NM_016650.1	NM_024021		NP_683876
219608_s_at	0.043703	gb:NM_024862.1 /DEF=Homo sapiens hypothetical protein FLJ13962 (FLJ13962), mRNA /FEA=mRNA /GEN=FLJ13962 /PROD=hypothetical protein FLJ13962 /DB_XREF=gi:13376291 /UG=Hs.330407 hypothetical protein FLJ13962 /FL=gb:NM_024862.1	NM_024862		NP_079138
219620_x_at	5.03E-04	gb:NM_017723.1 /DEF=Homo sapiens hypothetical protein FLJ20245 (FLJ20245), mRNA /FEA=mRNA /GEN=FLJ20245 /PROD=hypothetical protein FLJ20245 /DB_XREF=gi:8923220 /UG=Hs.169758 hypothetical protein FLJ20245 /FL=gb:NM_017723.1	NM_017723		NP_060193

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219625_s_at	0.003597	gb:NM_005713.1 /DEF=Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) binding protein (COL4A3BP), mRNA. /FEA=mRNA /GEN=COL4A3BP /PROD=collagen, type IV, alpha 3 (Goodpasture antigen)binding protein /DB_XREF=gi:5031716 /UG=Hs.21276 collagen, type IV, alpha 3 (Goodpasture antigen) binding protein. /FL=gb:AF136450.1 gb:NM_005713.1	NM_005713		NP_112729
219632_s_at	0.005121	gb:AF196175.1 /DEF=Homo sapiens capsaicin receptor mRNA, complete cds. /FEA=mRNA /PROD=capsaicin receptor /DB_XREF=gi:12003145 /UG=Hs.283010 vanilloid receptor subtype 1 /FL=gb:AF196175.1 gb:NM_018727.1	AF196175		NP_542437
219634_at	0.010205	gb:NM_018413.1 /DEF=Homo sapiens chondroitin 4-sulfotransferase (C4ST), mRNA. /FEA=mRNA /GEN=C4ST /PROD=chondroitin 4-sulfotransferase /DB_XREF=gi:8923757 /UG=Hs.287402 chondroitin 4-sulfotransferase /FL=gb:AB042326.1 gb:NM_018413.1 gb:AF239820.1	NM_018413		NP_060883
219639_x_at	0.004341	gb:NM_020213.1 /DEF=Homo sapiens hypothetical protein from EUROIMAGE 1977056 (LOC56965), mRNA. /FEA=mRNA /GEN=LOC56965 /PROD=hypothetical protein from EUROIMAGE 1977056 /DB_XREF=gi:9910373 /UG=Hs.315687 hypothetical protein from EUROIMAGE 1977056 /FL=gb:NM_020213.1	NM_020213		NP_064599

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219644_at	0.013088	gb:NM_016122.1 /DEF=Homo sapiens NY-REN-58 antigen (LOC51134), mRNA. /FEA=mRNA /GEN=LOC51134 /PROD=NY-REN-58 antigen /DB_XREF=gi:7705838 /UG=Hs.56148 NY-REN-58 antigen /FL=gb:AF155115.1 gb:NM_016122.1	NM_016122		NP_057206
219649_at	3.26E-04	gb:NM_013339.1 /DEF=Homo sapiens dolichyl-P-Glc:Man9GlcNAc2-PP-dolichylglucosyltransferase (ALG6), mRNA. /FEA=mRNA /GEN=ALG6 /PROD=dolichyl-P-Glc:Man9GlcNAc2-PP-dolichylglucosyltransferase /DB_XREF=gi:7019324 /UG=Hs.80042 dolichyl-P-Glc:Man9GlcNAc2-PP-dolichylglucosyltransferase /FL=gb:AF063604.1 gb:BC001253.1 gb:AF102851.1 gb:NM_013339.1	NM_013339		NP_037471
219657_s_at	0.013223	gb:NM_016531.1 /DEF=Homo sapiens hBKLF for basic kruppel like factor (LOC51274), mRNA. /FEA=mRNA /GEN=LOC51274 /PROD=hBKLF for basic kruppel like factor /DB_XREF=gi:7706076 /UG=Hs:119640 hBKLF for basic kruppel like factor /FL=gb:AF285837.1 gb:AB024523.1 gb:NM_016531.1	NM_016531		NP_057615
219666_at	1.26E-04	gb:NM_022349.1 /DEF=Homo sapiens CD20-like precursor (LOC64166), mRNA. /FEA=mRNA /GEN=LOC64166 /PROD=CD20-like precursor /DB_XREF=gi:11641258 /UG=Hs.17914 membrane-spanning 4-domains, subfamily A, member6 /FL=gb:AF142409.1 gb:NM_022349.1	NM_022349		NP_690591

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219667_s_at	0.007052	gb:NM_017935.1 /DEF=Homo sapiens hypothetical protein FLJ20706 (FLJ20706), mRNA. /FEA=mRNA /GEN=FLJ20706 /PROD=hypothetical protein FLJ20706 /DB_XREF=gi:8923635 /UG=Hs.193736 hypothetical protein FLJ20706 /FL=gb:NM_017935.1	NM_017935		NP_060405
219669_at	0.002536	gb:NM_020406.1 /DEF=Homo sapiens polycythemia rubra vera 1; cell surface receptor (PRV1), mRNA. /FEA=mRNA /GEN=PRV1 /PROD=polycythemia rubra vera 1; cell surfacereceptor /DB_XREF=gi:9966888 /UG=Hs.232165 polycythemia rubra vera 1; cell surface receptor /FL=gb:AF146747.1 gb:NM_020406.1	NM_020406		NP_065139
219672_at	0.008237	gb:NM_016633.1 /DEF=Homo sapiens EDRF protein (LOC51327), mRNA. /FEA=mRNA /GEN=LOC51327 /PROD=EDRF protein /DB_XREF=gi:7706179 /UG=Hs.274309 erythroid differentiation-related factor /FL=gb:AF208865.1 gb:NM_016633.1	NM_016633		NP_057717
219673_at	0.004341	gb:NM_017696.1 /DEF=Homo sapiens hypothetical protein FLJ20170 (FLJ20170), mRNA. /FEA=mRNA /GEN=FLJ20170 /PROD=hypothetical protein FLJ20170 /DB_XREF=gi:8923164 /UG=Hs.279008 hypothetical protein FLJ20170 /FL=gb:NM_017696.1	NM_017696		NP_060166
219680_at	0.041795	gb:NM_024618.1 /DEF=Homo sapiens hypothetical protein FLJ21478 (FLJ21478), mRNA. /FEA=mRNA /GEN=FLJ21478 /PROD=hypothetical protein FLJ21478 /DB_XREF=gi:13375837 /UG=Hs.31097 hypothetical protein FLJ21478 /FL=gb:NM_024618.1	NM_024618		NP_733840

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219694_at	1.26E-04	gb:NM_019018.1 /DEF=Homo sapiens hypothetical protein (FLJ11127); mRNA. /FEA=mRNA /GEN=FLJ11127 /PROD=hypothetical protein /DB_XREF=gi:9506640 /UG=Hs.91165 hypothetical protein /FL=gb:NM_019018.1	NM_019018		NP_061891
219700_at	3.56E-04	gb:NM_020405.1 /DEF=Homo sapiens tumor endothelial marker 7 precursor (TEM7); mRNA. /FEA=mRNA /GEN=TEM7 /PROD=tumor endothelial marker 7 precursor /DB_XREF=gi:9966886 /UG=Hs.125036 tumor endothelial marker 7 precursor /FL=gb:AF279144.1 gb:NM_020405.1	NM_020405		NP_065138
219711_at	0.026891	gb:NM_017652.1 /DEF=Homo sapiens hypothetical protein FLJ20070 (FLJ20070); mRNA. /FEA=mRNA /GEN=FLJ20070 /PROD=hypothetical protein FLJ20070 /DB_XREF=gi:8923076 /UG=Hs.186711 hypothetical protein FLJ20070 /FL=gb:NM_017652.1	NM_017652		NP_060122
219717_at	6.68E-04	gb:NM_017741.1 /DEF=Homo sapiens hypothetical protein FLJ20280 (FLJ20280); mRNA. /FEA=mRNA /GEN=FLJ20280 /PROD=hypothetical protein FLJ20280 /DB_XREF=gi:8923256 /UG=Hs.270134 hypothetical protein FLJ20280 /FL=gb:NM_017741.1	NM_017741		NP_060211
219734_at	0.00161	gb:NM_017699.1 /DEF=Homo sapiens hypothetical protein FLJ20174 (FLJ20174); mRNA. /FEA=mRNA /GEN=FLJ20174 /PROD=hypothetical protein FLJ20174 /DB_XREF=gi:8923170 /UG=Hs.114556 hypothetical protein FLJ20174 /FL=gb:NM_017699.1	NM_017699		NP_060169

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219770_at	0.005852	gb:NM_024659.1 /DEF=Homo sapiens hypothetical protein FLJ11753 (FLJ11753), mRNA. /FEA=mRNA /GEN=FLJ11753 /PROD=hypothetical protein FLJ11753 /DB_XREF=gi:13375910 /UG=Hs.62348 hypothetical protein FLJ11753 /FL=gb:NM_024659.1	NM_024659		NP_078935
219797_at	7.58E-04	gb:NM_012214.1 /DEF=Homo sapiens mannosyl (alpha-1,3)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme A (MGAT4A), mRNA. /FEA=mRNA /GEN=MGAT4A /PROD=mannosyl (alpha-1,3)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme A /DB_XREF=gi:6912501 /UG=Hs.177576 mannosyl (alpha-1,3)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme A /FL=gb:AB000616.1 gb:NM_012214.1	NM_012214		NP_036346
219799_s_at	0.033381	gb:NM_005771.1 /DEF=Homo sapiens retinol dehydrogenase homolog (RDHL), mRNA. /FEA=mRNA /GEN=RDHL /PROD=retinol dehydrogenase homolog /DB_XREF=gi:5032034 /UG=Hs.179608 retinol dehydrogenase homolog /FL=gb:AF067174.1 gb:NM_005771.1	NM_005771		NP_005762
219806_s_at	1.26E-04	gb:NM_020179.1 /DEF=Homo sapiens FN5 protein (FN5), mRNA. /FEA=mRNA /GEN=FN5 /PROD=FN5 protein /DB_XREF=gi:9910225 /UG=Hs.259737 FN5 protein /FL=gb:AF197137.1 gb:NM_020179.1	NM_020179		NP_064564

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219812_at	6.68E-04	gb:NM_024070.1 /DEF=Homo sapiens hypothetical protein MGC2463 (MGC2463), mRNA. /FEA=mRNA /GEN=MGC2463 /PROD=hypothetical protein MGC2463 /DB_XREF=gi:13129051 /UG=Hs.323634 hypothetical protein MGC2463 /FL=gb:BC001129.1 gb:NM_024070.1	NM_024070		
219816_s_at	3.26E-04	gb:NM_018107.1 /DEF=Homo sapiens hypothetical protein FLJ10482 (FLJ10482), mRNA. /FEA=mRNA /GEN=FLJ10482 /PROD=hypothetical protein FLJ10482 /DB_XREF=gi:8922449 /UG=Hs.4997 hypothetical protein FLJ10482 /FL=gb:NM_018107.1	NM_018107		NP_060577
219846_at	0.016934	gb:NM_025174.1 /DEF=Homo sapiens hypothetical protein FLJ23040 (FLJ23040), mRNA. /FEA=mRNA /GEN=FLJ23040 /PROD=hypothetical protein FLJ23040 /DB_XREF=gi:13376769 /UG=Hs.169813 hypothetical protein FLJ23040 /FL=gb:NM_025174.1	NM_025174		NP_079450
219859_at	1.26E-04	gb:NM_014358.1 /DEF=Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9 (CLECSF9), mRNA. /FEA=mRNA /GEN=CLECSF9 /PROD=C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9 /DB_XREF=gi:7657332 /UG=Hs.236516 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9 /FL=gb:BC000715.1 gb:AB024718.1 gb:NM_014358.1	NM_014358		NP_055173

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
219860_at	0.010205	gb:NM_025262.1 /DEF=Homo sapiens G5C protein (G5C), mRNA. /FEA=mRNA /GEN=G5C /PROD=G5C protein /DB_XREF=gi:13376875 /UG=Hs.246845 G5C protein /FL=gb:NM_025262.1	NM_025262		NP_079538
219862_s_at	0.007779	gb:NM_012336.1 /DEF=Homo sapiens nuclear prelamin A recognition factor (NARF), mRNA. /FEA=mRNA /GEN=NARF /PROD=nuclear prelamin A recognition factor /DB_XREF=gi:6912523 /UG=Hs.256526 nuclear prelamin A recognition factor /FL=gb:AF128406.1 gb:NM_012336.1	NM_012336		NP_114174
219868_s_at	0.039278	gb:NM_016376.1 /DEF=Homo sapiens ANKHZN protein (ANKHZN), mRNA. /FEA=mRNA /GEN=ANKHZN /PROD=ANKHZN protein /DB_XREF=gi:7705277 /UG=Hs.284163 ANKHZN protein /FL=gb:AB037360.1 gb:NM_016376.1	NM_016376		NP_065791
219869_s_at	9.79E-04	gb:NM_022154.1 /DEF=Homo sapiens up-regulated by BCG-CWS (LOC64116), mRNA. /FEA=mRNA /GEN=LOC64116 /PROD=up-regulated by BCG-CWS /DB_XREF=gi:11545899 /UG=Hs.284205 up-regulated by BCG-CWS /FL=gb:NM_022154.1	NM_022154		NP_071437
219885_at	0.046329	gb:NM_018042.1 /DEF=Homo sapiens hypothetical protein FLJ10260 (FLJ10260), mRNA. /FEA=mRNA /GEN=FLJ10260 /PROD=hypothetical protein FLJ10260 /DB_XREF=gi:8922317 /UG=Hs.105859 hypothetical protein FLJ10260 /FL=gb:NM_018042.1	NM_018042		NP_060512

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
219892_at	1.26E-04	gb:NM_023003.1 /DEF=Homo sapiens transmembrane 6 superfamily member 1 (TM6SF1), mRNA. /FEA=mRNA /GEN=TM6SF1 /PROD=transmembrane 6 superfamily member 1 /DB_XREF=gi:13194198 /UG=Hs.133865. transmembrane. 6 superfamily member 1 /FL=gb:AF255922.1 gb:NM_023003.1	NM_023003		NP_075379
219929_s_at	8.34E-04	gb:NM_024071.1 /DEF=Homo sapiens hypothetical protein MGC2550 (MGC2550), mRNA. /FEA=mRNA /GEN=MGC2550 /PROD=hypothetical protein MGC2550 /DB_XREF=gi:13129053 /UG=Hs.318498 hypothetical protein. MGC2550 /FL=gb:BC001130.1 gb:NM_024071.1	NM_024071		NP_076976
219931_s_at	0.002536	gb:NM_021633.1 /DEF=Homo sapiens kelch-like protein C3IP1 (C3IP1), mRNA. /FEA=mRNA /GEN=C3IP1 /PROD=kelch-like protein C3IP1 /DB_XREF=gi:11056005 /UG=Hs.3826 kelch-like protein C3IP1 /FL=gb:AF190900.1 gb:NM_021633.1 gb:BC003183.1 gb:BC004175.1	NM_021633		NP_067646
219933_at	0.016934	gb:NM_016066.1 /DEF=Homo sapiens CGI-133 protein (LOC51022), mRNA. /FEA=mRNA /GEN=LOC51022 /PROD=CGI-133 protein /DB_XREF=gi:7705627 /UG=Hs.5054 CGI-133 protein /FL=gb:AF151891.1 gb:NM_016066.1 gb:AF132495.2	NM_016066		NP_057150

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219938_s_at	3.26E-04	gb:NM_024430.1 /DEF=Homo sapiens proline-serine-threonine phosphatase interacting protein 2 (PSTPIP2), mRNA. /FEA=mRNA /GEN=PSTPIP2 /PROD=proline-serine-threonine phosphatase interacting protein 2 /DB_XREF=gi:13270468 /UG=Hs.69149 proline-serine-threonine phosphatase interacting protein 2 /FL=gb:NM_024430.1	NM_024430		NP_077748
219947_at	0.001116	gb:NM_016184.1 /DEF=Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 (CLECSF6), mRNA. /FEA=mRNA /GEN=CLECSF6 /PROD=C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 /DB_XREF=gi:7705337 /UG=Hs.115515 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 /FL=gb:AF109146.1 gb:NM_016184.1 gb:AF067800.1	NM_016184		NP_057268
219952_s_at	0.007934	gb:NM_020533.1 /DEF=Homo sapiens mucolipin 1 (MCOLN1), mRNA. /FEA=mRNA /GEN=MCOLN1 /PROD=mucolipin 1 /DB_XREF=gi:10092596 /UG=Hs.12909 mucolipin 1 /FL=gb:NM_020533.1 gb:BC005149.1 gb:AF287269.1 gb:AF249319.1	NM_020533		NP_065394

Gene Identifier	p-value	Description	Gen Accession No.	Unig ne Accession No.	Protein Accession No.
219960_s_at	0.010205	gb:NM_015984.1 /DEF=Homo sapiens ubiquitin C-terminal hydrolase UCH37 (UCH37), mRNA. /FEA=mRNA /GEN=UCH37 /PROD=ubiquitin C-terminal hydrolase UCH37 /DB_XREF=gi:7706752 /UG=Hs.171581 ubiquitin C-terminal hydrolase UCH37 /FL=gb:AF147717.1 gb:NM_015984.1	NM_015984		NP_057068
219963_at	0.004659	gb:NM_016364.1 /DEF=Homo sapiens protein phosphatase (LOC51207), mRNA. /FEA=mRNA /GEN=LOC51207 /PROD=protein phosphatase /DB_XREF=gi:7705958 /UG=Hs.178170 protein phosphatase /FL=gb:AB027004.1 gb:NM_016364.1	NM_016364		NP_057448
219974_x_at	0.02145	gb:NM_018479.1 /DEF=Homo sapiens uncharacterized hypothalamus protein HCDASE (LOC55862), mRNA. /FEA=mRNA /GEN=LOC55862 /PROD=uncharacterized hypothalamus protein HCDASE /DB_XREF=gi:8923863 /UG=Hs.239218 uncharacterized hypothalamus protein HCDASE /FL=gb:AF220192.1 gb:NM_018479.1	NM_018479		NP_060949
219979_s_at	0.003171	gb:NM_016401.1 /DEF=Homo sapiens hypothetical protein (HSPC138), mRNA. /FEA=mRNA /GEN=HSPC138 /PROD=hypothetical protein /DB_XREF=gi:7705470 /UG=Hs.283775 hypothetical protein /FL=gb:AF161487.1 gb:NM_016401.1	NM_016401		NP_057485
219981_x_at	0.001116	gb:NM_017961.1 /DEF=Homo sapiens hypothetical protein FLJ20813 (FLJ20813), mRNA. /FEA=mRNA /GEN=FLJ20813 /PROD=hypothetical protein FLJ20813 /DB_XREF=gi:8923685 /UG=Hs.306203 hypothetical protein FLJ20813 /FL=gb:NM_017961.1	NM_017961		NP_060431

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
219994_at	0.028893	gb:NM_019043.1 /DEF=Homo sapiens similar to proline-rich protein 48 (LOC54518), mRNA. /FEA=mRNA /GEN=LOC54518 /PROD=similar to proline-rich protein 48 /DB_XREF=gi:9506856 /UG=Hs.98874 similar to proline-rich protein 48 /FL=gb:NM_019043.1	NM_019043		NP_061916
220000_at	3.09E-04	gb:NM_003830.1 /DEF=Homo sapiens sialic acid binding Ig-like lectin 5 (SIGLEC5), mRNA. /FEA=mRNA /GEN=SIGLEC5 /PROD=sialic acid binding Ig-like lectin 5 /DB_XREF=gi:4502658 /UG=Hs.117005 sialic acid binding Ig-like lectin 5 /FL=gb:U71383.1 gb:NM_003830.1 gb:AF170484.1	NM_003830		NP_003821
220001_at	0.003171	gb:NM_012387.1 /DEF=Homo sapiens peptidyl arginine deiminase, type V (PAD), mRNA. /FEA=mRNA /GEN=PAD /PROD=peptidyl arginine deiminase, type V /DB_XREF=gi:6912575 /UG=Hs.117232 peptidyl arginine deiminase, type V /FL=gb:AB017919.1 gb:NM_012387.1	NM_012387		NP_036519
220005_at	1.26E-04	gb:NM_023914.1 /DEF=Homo sapiens G protein-coupled receptor 86 (GPR86), mRNA. /FEA=mRNA /GEN=GPR86 /PROD=G protein-coupled receptor 86 /DB_XREF=gi:13194202 /UG=Hs.13040 G protein-coupled receptor 86 /FL=gb:AF295368.1 gb:NM_023914.1 gb:AF178982.1 gb:AF345565.1	NM_023914		NP_795713

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220009_at	0.008237	gb:NM_024778.1 /DEF=Homo sapiens hypothetical protein FLJ22612 (FLJ22612), mRNA. /FEA=mRNA /GEN=FLJ22612 /PROD=hypothetical protein FLJ22612 /DB_XREF=gi:13376133 /UG=Hs.144266 hypothetical protein FLJ22612 /FL=gb:NM_024778.1	NM_024778		NP_079054
220015_at	0.004341	gb:NM_017766.1 /DEF=Homo sapiens hypothetical protein FLJ20321 (FLJ20321), mRNA. /FEA=mRNA /GEN=FLJ20321 /PROD=hypothetical protein FLJ20321 /DB_XREF=gi:8923302 /UG=Hs.162196 hypothetical protein FLJ20321 /FL=gb:BC004410.1 gb:NM_017766.1	NM_017766		NP_060236
220023_at	0.016934	gb:NM_018690.1 /DEF=Homo sapiens apolipoprotein B48 receptor (APOB48R), mRNA. /FEA=mRNA /GEN=APOB48R /PROD=apolipoprotein B48 receptor /DB_XREF=gi:8922078 /UG=Hs.200333 apolipoprotein B48 receptor /FL=gb:AF141332.1 gb:AF141334.1 gb:NM_018690.1	NM_018690		NP_061160
220034_at	0.003522	gb:NM_007199.1 /DEF=Homo sapiens interleukin-1 receptor-associated kinase M (IRAK-M), mRNA. /FEA=mRNA /GEN=IRAK-M /PROD=interleukin-1 receptor-associated kinase M /DB_XREF=gi:6005791 /UG=Hs.268552 interleukin-1 receptor-associated kinase M /FL=gb:AF113136.1 gb:NM_007199.1	NM_007199		NP_009130

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220050_at	0.00423	gb:NM_018956.1 /DEF=Homo sapiens chromosome 9 open reading frame 9 (C9ORF9), mRNA. /FEA=mRNA /GEN=C9ORF9 /PROD=chromosome 9 open reading frame 9 /DB_XREF=gi:9506440 /UG=Hs.62595 chromosome 9 open reading frame 9 /FL=gb:NM_018956.1	NM_018956		NP_061829
220052_s_at	0.00161	gb:NM_012461.1 /DEF=Homo sapiens TERF1 (TRF1)-interacting nuclear factor 2 (TINF2), mRNA. /FEA=mRNA /GEN=TINF2 /PROD=TERF1 (TRF1)-interacting nuclear factor 2 /DB_XREF=gi:6912715 /UG=Hs.7797 TERF1 (TRF1)-interacting nuclear factor 2 /FL=gb:AF195512.1 gb:NM_012461.1	NM_012461		NP_036593
220071_x_at	0.00228	gb:NM_018097.1 /DEF=Homo sapiens hypothetical protein FLJ10460 (FLJ10460), mRNA. /FEA=mRNA /GEN=FLJ10460 /PROD=hypothetical protein FLJ10460 /DB_XREF=gi:8922429 /UG=Hs.14347 hypothetical protein FLJ10460 /FL=gb:NM_018097.1	NM_018097		NP_060567
220078_at	0.001618	gb:NM_018391.1 /DEF=Homo sapiens hypothetical protein FLJ11328 (FLJ11328), mRNA. /FEA=mRNA /GEN=FLJ11328 /PROD=hypothetical protein FLJ11328 /DB_XREF=gi:8922996 /UG=Hs.178518 hypothetical protein FLJ11328 /FL=gb:NM_018391.1	NM_018391		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220088_at	0.001116	gb:NM_001736.1 /DEF=Homo sapiens complement component 5 receptor 1 (C5a ligand) (C5R1), mRNA. /FEA=mRNA. /GEN=C5R1 /PROD=complement component 5 receptor 1 (C5a ligand) /DB_XREF=gi:4502508 /UG=Hs.2161 complement component 5 receptor 1 (C5a ligand) /FL=gb:M62505.1 gb:NM_001736.1	NM_001736		NP_001727
220097_s_at	0.046435	gb:NM_017728.1 /DEF=Homo sapiens hypothetical protein FLJ20255 (FLJ20255), mRNA. /FEA=mRNA /GEN=FLJ20255 /PROD=hypothetical protein FLJ20255 /DB_XREF=gi:8923229 /UG=Hs.274247 hypothetical protein FLJ20255 /FL=gb:NM_017728.1	NM_017728		NP_060198
220112_at	0.00161	gb:NM_024669.1 /DEF=Homo sapiens hypothetical protein FLJ11795 (FLJ11795), mRNA. /FEA=mRNA /GEN=FLJ11795 /PROD=hypothetical protein FLJ11795 /DB_XREF=gi:13375927 /UG=Hs.84560 hypothetical protein FLJ11795 /FL=gb:NM_024669.1	NM_024669		NP_078945
220113_x_at	0.001116	gb:NM_019014.1 /DEF=Homo sapiens similar to DNA-directed RNA polymerase I (135 kDa) (FLJ10816), mRNA. /FEA=mRNA /GEN=FLJ10816 /PROD=similar to DNA-directed RNA polymerase I (135kDa) /DB_XREF=gi:9506618 /UG=Hs.86337 similar to DNA-directed RNA polymerase I (135 kDa) /FL=gb:NM_019014.1	NM_019014		NP_115588

Gene Identifier	p-value	D scription	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220122_at	0.033381	gb:NM_024717.1 /DEF=Homo sapiens hypothetical protein FLJ22344 (FLJ22344), mRNA. /FEA=mRNA /GEN=FLJ22344 /PROD=hypothetical protein FLJ22344 /DB_XREF=gi:13376020 /UG=Hs.107716 hypothetical protein. FLJ22344 /FL=gb:NM_024717.1	NM_024717		NP_078993
220127_s_at	0.041055	gb:NM_017703.1 /DEF=Homo sapiens hypothetical protein FLJ20188 (FLJ20188), mRNA. /FEA=mRNA /GEN=FLJ20188 /PROD=hypothetical protein FLJ20188 /DB_XREF=gi:8923178 /UG=Hs.12439 hypothetical protein FLJ20188 /FL=gb:NM_017703.1	NM_017703		NP_060173
220132_s_at	0.016934	gb:NM_013269.1 /DEF=Homo sapiens lectin-like NK cell receptor (LLT1), mRNA. /FEA=mRNA /GEN=LLT1 /PROD=lectin-like NK cell receptor /DB_XREF=gi:7019446 /UG=Hs.136748 lectin-like NK cell receptor /FL=gb:AF133299.1 gb:NM_013269.1 gb:AF285087.1	NM_013269		NP_037401
220138_at	5.07E-04	gb:NM_004821.1 /DEF=Homo sapiens heart and neural crest derivatives expressed 1 (HAND1); mRNA. /FEA=mRNA /GEN=HAND1 /PROD=basic helix-loop-helix transcription factorHAND1 /DB_XREF=gi:4758505 /UG=Hs.152531 heart and neural crest derivatives expressed 1 /FL=gb:AF061756.1 gb:NM_004821.1	NM_004821		NP_004812
220140_s_at	0.033381	gb:NM_013323.1 /DEF=Homo sapiens sorting nexin 11 (SNX11), mRNA. /FEA=mRNA /GEN=SNX11 /PROD=sorting nexin 11 /DB_XREF=gi:7019538 /UG=Hs.15827 sorting nexin 11 /FL=gb:AF121861.1 gb:NM_013323.1	NM_013323		NP_689450

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
220147_s_at	0.013223	gb:NM_021238.1 /DEF=Homo sapiens TERA protein (TERA), mRNA. /FEA=mRNA /GEN=TERA /PROD=TERA protein /DB_XREF=gi:10864048 /UG=Hs.180780 TERA protein /FL=gb:NM_021238.1 gb:BC000024.1	NM_021238		NP_067061
220162_s_at	0.001012	gb:NM_022352.1 /DEF=Homo sapiens caspase recruitment domain protein 9 (LOC64170), mRNA. /FEA=mRNA /GEN=LOC64170 /PROD=caspase recruitment domain protein 9 /DB_XREF=gi:11641262 /UG=Hs.271815 caspase recruitment domain protein 9 /FL=gb:NM_022352.1	NM_022352		NP_434701
220175_s_at	0.041055	gb:NM_020667.1 /DEF=Homo sapiens hypothetical protein from clone 1659351 (LOC57397), mRNA. /FEA=mRNA /GEN=LOC57397 /PROD=hypothetical protein from clone 1659351 /DB_XREF=gi:10190707 /UG=Hs.288838 hypothetical protein from clone 1659351 /FL=gb:NM_020667.1	NM_020667		
220202_s_at	0.00228	gb:NM_018835.1 /DEF=Homo sapiens membrane-associated nucleic acid binding protein (MNAB), mRNA. /FEA=mRNA /GEN=MNAB /PROD=membrane-associated nucleic acid binding protein /DB_XREF=gi:9256536 /UG=Hs.112227 membrane-associated nucleic acid binding protein /FL=gb:NM_018835.1	NM_018835		NP_061323

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220232_at	0.042536	gb:NM_024906.1 /DEF=Homo sapiens hypothetical protein FLJ21032 (FLJ21032), mRNA. /FEA=mRNA /GEN=FLJ21032 /PROD=hypothetical protein FLJ21032 /DB_XREF=gi:13376362 /UG=Hs.247474 hypothetical protein FLJ21032 /FL=gb:NM_024906.1	NM_024906		NP_079182
220235_s_at	1.26E-04	gb:NM_018372.1 /DEF=Homo sapiens hypothetical protein FLJ11269 (FLJ11269), mRNA. /FEA=mRNA /GEN=FLJ11269 /PROD=hypothetical protein FLJ11269 /DB_XREF=gi:8922961 /UG=Hs.25245 hypothetical protein FLJ11269 /FL=gb:NM_018372.1	NM_018372		NP_060842
220244_at	0.007779	gb:NM_013343.1 /DEF=Homo sapiens NAG-7 protein (NAG-7), mRNA. /FEA=mRNA /GEN=NAG-7 /PROD=NAG-7 protein /DB_XREF=gi:7106370 /UG=Hs.278951 NAG-7 protein /FL=gb:AF086709.2 gb:NM_013343.1	NM_013343		NP_037475
220292_at	0.002464	gb:NM_024340.1 /DEF=Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA. /FEA=mRNA /GEN=MGC4179 /PROD=hypothetical protein MGC4179 /DB_XREF=gi:13236590 /UG=Hs.129369 hypothetical protein MGC4179 /FL=gb:BC002859.1 gb:NM_024340.1	NM_024340		
220305_at	3.26E-04	gb:NM_024030.1 /DEF=Homo sapiens hypothetical protein MGC3260 (MGC3260), mRNA. /FEA=mRNA /GEN=MGC3260 /PROD=hypothetical protein MGC3260 /DB_XREF=gi:13128977 /UG=Hs.15514 hypothetical protein MGC3260 /FL=gb:BC000073.1 gb:NM_024030.1	NM_024030		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220306_at	0.004026	gb:NM_017709.1 /DEF=Homo sapiens hypothetical protein FLJ20202 (FLJ20202), mRNA. /FEA=mRNA /GEN=FLJ20202 /PROD=hypothetical protein FLJ20202 /DB_XREF=gi:8923191 /UG=Hs.155556 hypothetical protein FLJ20202 /FL=gb:NM_017709.1	NM_017709		NP_060179
220330_s_at	1.26E-04	gb:NM_022136.1 /DEF=Homo sapiens SAM domain, SH3 domain and nuclear localisation signals, 1 (SAMSN1), mRNA. /FEA=mRNA /GEN=SAMSN1 /PROD=SAM domain, SH3 domain and nuclear localisationsignals, 1 /DB_XREF=gi:11545870 /UG=Hs.24633 SAM domain, SH3 domain and nuclear localisation signals, 1 /FL=gb:AF222927.1 gb:NM_022136.1	NM_022136		NP_071419
220363_s_at	0.004452	gb:NM_022086.1 /DEF=Homo sapiens hypothetical protein FLJ11656 (FLJ11656), mRNA. /FEA=mRNA /GEN=FLJ11656 /PROD=hypothetical protein FLJ11656 /DB_XREF=gi:11545798 /UG=Hs.96560 hypothetical protein FLJ11656 /FL=gb:NM_022086.1	NM_022086		NP_573403
220367_s_at	2.05E-04	gb:NM_024545.1 /DEF=Homo sapiens hypothetical protein FLJ12761 (FLJ12761), mRNA. /FEA=mRNA /GEN=FLJ12761 /PROD=hypothetical protein FLJ12761 /DB_XREF=gi:13375706 /UG=Hs.10554 hypothetical protein FLJ12761 /FL=gb:NM_024545.1	NM_024545		NP_078821

Gen Id ntifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Acc ssion No.
220368_s_at	0.00228	gb:NM_017936.1 /DEF=Homo sapiens hypothetical protein FLJ20707 (FLJ20707), mRNA. /FEA=mRNA /GEN=FLJ20707 /PROD=hypothetical protein FLJ20707 /DB_XREF=gi:8923637 /UG=Hs.109441 hypothetical protein FLJ20707 /FL=gb:NM_017936.1	NM_017936		NP_115949
220375_s_at	0.035763	gb:NM_024752.1 /DEF=Homo sapiens hypothetical protein FLJ23312 (FLJ23312), mRNA. /FEA=mRNA /GEN=FLJ23312 /PROD=hypothetical protein FLJ23312 /DB_XREF=gi:13376081 /UG=Hs.126555 hypothetical protein FLJ23312 /FL=gb:NM_024752.1	NM_024752		
220404_at	0.010205	gb:NM_014076.1 /DEF=Homo sapiens PRO0611 protein (PRO0611), mRNA. /FEA=mRNA /GEN=PRO0611 /PROD=PRO0611 protein /DB_XREF=gi:7662571 /UG=Hs.163833 PRO0611 protein /FL=gb:AF111851.1 gb:NM_014076.1	NM_014076		
220416_at	1.26E-04	gb:NM_024837.1 /DEF=Homo sapiens hypothetical protein FLJ21472 (FLJ21472), mRNA. /FEA=mRNA /GEN=FLJ21472 /PROD=hypothetical protein FLJ21472 /DB_XREF=gi:13376248 /UG=Hs.182738 hypothetical protein FLJ21472 /FL=gb:NM_024837.1	NM_024837		NP_079113
220417_s_at	0.013223	gb:NM_015963.1 /DEF=Homo sapiens CGI-36 protein (LOC51078), mRNA. /FEA=mRNA /GEN=LOC51078 /PROD=CGI-36 protein /DB_XREF=gi:7705731 /UG=Hs.182986 CGI-36 protein /FL=gb:AF132970.1 gb:NM_015963.1	NM_015963		NP_057047

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220477_s_at	0.034221	gb:NM_014145.1 /DEF=Homo sapiens HSPC274 protein (HSPC274), mRNA. /FEA=mRNA /GEN=HSPC274 /PROD=HSPC274 protein /DB_XREF=gi:7661833 /UG=Hs.323476 HSPC274 protein /FL=gb:AF161392.1 gb:NM_014145.1	NM_014145		NP_054864
220491_at	0.014208	gb:NM_021175.1 /DEF=Homo sapiens liver-expressed antimicrobial peptide (LEAP-1); mRNA. /FEA=mRNA /GEN=LEAP-1 /PROD=liver-expressed antimicrobial peptide /DB_XREF=gi:10863972 /UG=Hs.8821 liver-expressed antimicrobial peptide /FL=gb:AF309489.1 gb:NM_021175.1 gb:AF131292.1	NM_021175		NP_066998
220494_s_at	0.043705	gb:NM_018678.1 /DEF=Homo sapiens lipopolysaccharide specific response-68 protein (LSR68), mRNA. /FEA=mRNA /GEN=LSR68 /PROD=lipopolysaccharide specific response-68 protein /DB_XREF=gi:8923914 /UG=Hs.103189 lipopolysaccharide specific response-68 protein /FL=gb:AF147723.1 gb:NM_018678.1	NM_018678		NP_061148
220500_s_at	0.009275	gb:NM_007082.2 /DEF=Homo sapiens RAB, member of RAS oncogene family-like 2A (RABL2A), mRNA. /FEA=mRNA /GEN=RABL2A /PROD=RAB, member of RAS oncogene family-like 2A /DB_XREF=gi:7549818 /UG=Hs.122660 RAB, member of RAS oncogene family-like 2A /FL=gb:AF095350.1 gb:NM_007082.2	NM_007082		NP_038198

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220532_s_at	0.013223	gb:NM_014020.1 /DEF=Homo sapiens LR8 protein (LR8), mRNA. /FEA=mRNA /GEN=LR8 /PROD=LR8 protein /DB_XREF=gi:7662497. /UG=Hs.190161 LR8 protein /FL=gb:AF115384.1 gb:NM_014020.1	NM_014020		NP_054739
220537_at	0.007063	gb:NM_017677.1 /DEF=Homo sapiens hypothetical protein FLJ20126 (FLJ20126), mRNA. /FEA=mRNA /GEN=FLJ20126 /PROD=hypothetical protein FLJ20126. /DB_XREF=gi:8923125 /UG=Hs.196568 hypothetical protein FLJ20126 /FL=gb:NM_017677.1	NM_017677		NP_060147
220570_at	0.041795	gb:NM_020415.2 /DEF=Homo sapiens found in inflammatory zone 3 (FIZZ3); mRNA. /FEA=mRNA /GEN=FIZZ3 /PROD=resistin. /DB_XREF=gi:13435379 /UG=Hs.283091 found in inflammatory zone, 3 /FL=gb:AF323081.1 gb:NM_020415.2 gb:AF205952.1	NM_020415		NP_065148
220577_at	1.26E-04	gb:NM_025006.1 /DEF=Homo sapiens hypothetical protein FLJ13373 (FLJ13373); mRNA. /FEA=mRNA /GEN=FLJ13373 /PROD=hypothetical protein FLJ13373 /DB_XREF=gi:13376522 /UG=Hs.287567 hypothetical protein FLJ13373 /FL=gb:NM_025006.1	NM_025006		NP_079282
220591_s_at	0.005852	gb:NM_025184.1 /DEF=Homo sapiens hypothetical protein FLJ22843 (FLJ22843), mRNA. /FEA=mRNA /GEN=FLJ22843 /PROD=hypothetical protein FLJ22843 /DB_XREF=gi:13376775 /UG=Hs.301143 hypothetical protein FLJ22843 /FL=gb:NM_025184.1	NM_025184		NP_079460

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
220597_s_at	0.02145	gb:NM_018694.1 /DEF=Homo sapiens HSVI binding protein (LOC55913), mRNA. /FEA=mRNA /GEN=LOC55913 /PROD=HSV1 binding protein /DB_XREF=gi:8923906 /UG=Hs.306208 HSVI binding protein. /FL=gb:AF267748.1 gb:NM_018694.1	NM_018694		NP_061164
220603_s_at	2.05E-04	gb:NM_018349.1 /DEF=Homo sapiens hypothetical protein FLJ11175 (FLJ11175), mRNA. /FEA=mRNA /GEN=FLJ11175 /PROD=hypothetical protein FLJ11175 /DB_XREF=gi:8922916 /UG=Hs.33368 hypothetical protein FLJ11175 /FL=gb:NM_018349.1	NM_018349		NP_060819
220610_s_at	0.029713	gb:NM_006309.1 /DEF=Homo sapiens leucine rich repeat (in FLII) interacting protein 2 (LRRFIP2), mRNA. /FEA=mRNA /GEN=LRRFIP2 /PROD=leucine rich repeat (in FLII) interacting protein 2 /DB_XREF=gi:5453725 /UG=Hs.57672 leucine rich repeat (in FLII) interacting protein 2 /FL=gb:AF115509.1 gb:NM_006309.1	NM_006309		NP_060194
220615_s_at	2.05E-04	gb:NM_018099.1 /DEF=Homo sapiens hypothetical protein FLJ10462 (FLJ10462), mRNA. /FEA=mRNA /GEN=FLJ10462 /PROD=hypothetical protein FLJ10462 /DB_XREF=gi:8922433 /UG=Hs.100895 hypothetical protein FLJ10462 /FL=gb:AL136843.1 gb:NM_018099.1	NM_018099		NP_060569
220661_s_at	1.26E-04	gb:NM_017865.1 /DEF=Homo sapiens hypothetical protein FLJ20531 (FLJ20531), mRNA. /FEA=mRNA /GEN=FLJ20531 /PROD=hypothetical protein FLJ20531 /DB_XREF=gi:8923497 /UG=Hs.23617 hypothetical protein FLJ20531 /FL=gb:NM_017865.1	NM_017865		NP_060335

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220669_at	0.041795	gb:NM_017493.2 /DEF=Homo sapiens Hin-1 (HSHIN1), mRNA. /FEA=mRNA /GEN=HSHIN1. /PROD=Hin-1 /DB_XREF=gi:13518029 /UG=Hs.252722 Hin-1 /FL=gb:NM_017493.2	NM_017493		NP_059963
220684_at	0.003171	gb:NM_013351.1 /DEF=Homo sapiens T-box 21 (TBX21), mRNA. /FEA=mRNA /GEN=TBX21 /PROD=T-box 21 /DB_XREF=gi:7019548 /UG=Hs.272409 T-box 21 /FL=gb:AF093098.1 gb:NM_013351.1 gb:AF241243.2	NM_013351		NP_037483
220691_at	0.026891	gb:NM_014114.1 /DEF=Homo sapiens PRO0097 protein (PRO0097), mRNA. /FEA=mRNA /GEN=PRO0097 /PROD=PRO0097 protein /DB_XREF=gi:7662521 /UG=Hs.278928 PRO0097 protein /FL=gb:AF090890.1 gb:NM_014114.1	NM_014114		NP_054833
220710_at	0.007258	gb:NM_024970.1 /DEF=Homo sapiens hypothetical protein FLJ11722 (FLJ11722), mRNA. /FEA=mRNA /GEN=FLJ11722 /PROD=hypothetical protein FLJ11722 /DB_XREF=gi:13376465 /UG=Hs.287444 hypothetical protein FLJ11722 /FL=gb:NM_024970.1	NM_024970		NP_079246
220739_s_at	1.26E-04	gb:NM_017623.1 /DEF=Homo sapiens hypothetical protein FLJ20018 (FLJ20018), mRNA. /FEA=mRNA /GEN=FLJ20018 /PROD=hypothetical protein FLJ20018 /DB_XREF=gi:8923022 /UG=Hs.44095 hypothetical protein FLJ20018 /FL=gb:NM_017623.1	NM_017623		NP_060093

Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
220741_s_at	1.26E-04	gb:NM_006903.1 /DEF=Homo sapiens inorganic pyrophosphatase (SID6-306), mRNA. /FEA=mRNA /GEN=SID6-306 /PROD=inorganic pyrophosphatase /DB_XREF=gi:13491177 /UG=Hs.5123 inorganic pyrophosphatase /FL=gb:NM_006903.1 gb:AB026722.1	NM_006903		NP_789845
220746_s_at	0.041055	gb:NM_016290.1 /DEF=Homo sapiens retinoid x receptor interacting protein (LOC51720), mRNA. /FEA=mRNA /GEN=LOC51720 /PROD=retinoid x receptor interacting protein /DB_XREF=gi:7706482 /UG=Hs.7889 retinoid x receptor interacting protein /FL=gb:AF113538.1 gb:NM_016290.1	NM_016290		NP_057374
220750_s_at	0.013223	gb:NM_022356.1 /DEF=Homo sapiens growth suppressor 1 (GROS1), mRNA. /FEA=mRNA /GEN=GROS1 /PROD=growth suppressor 1 /DB_XREF=gi:11641270 /UG=Hs.10114 growth suppressor 1 /FL=gb:AF097431.1 gb:NM_022356.1	NM_022356		NP_071751
220753_s_at	0.048741	gb:NM_015974.1 /DEF=Homo sapiens lambda-crystallin (LOC51084), mRNA. /FEA=mRNA /GEN=LOC51084 /PROD=lambda-crystallin /DB_XREF=gi:7705743 /UG=Hs.108896 lambda-crystallin /FL=gb:AF077049.1 gb:NM_015974.1	NM_015974		NP_057058

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
220757_s_at	0.02145	gb:NM_025241.1 /DEF=Homo sapiens UBX domain-containing gene 1 (UBXD1), mRNA. /FEA=mRNA /GEN=UBXD1 /PROD=UBX domain-containing gene 1 /DB_XREF=gi:13376853 /UG=Hs.11081 UBX domain-containing 2 /FL=gb:AF272893.1 gb:NM_025241.1	NM_025241		NP_079517
220761_s_at	0.00161	gb:NM_016281.1 /DEF=Homo sapiens STE20-like kinase (JIK), mRNA. /FEA=mRNA /GEN=JIK /PROD=STE20-like kinase /DB_XREF=gi:7705559 /UG=Hs.12040 STE20-like kinase /FL=gb:AF179867.1 gb:NM_016281.1	NM_016281		NP_057365
220796_x_at	0.033381	gb:NM_024881.1 /DEF=Homo sapiens hypothetical protein FLJ14251 (FLJ14251), mRNA. /FEA=mRNA /GEN=FLJ14251 /PROD=hypothetical protein FLJ14251 /DB_XREF=gi:13376323 /UG=Hs.214178 hypothetical protein FLJ14251 /FL=gb:NM_024881.1	NM_024881		NP_079157
220800_s_at	0.018693	gb:NM_014547.1 /DEF=Homo sapiens tropomodulin 3 (ubiquitous) (TMOD3), mRNA. /FEA=mRNA /GEN=TMOD3 /PROD=tropomodulin 3 (ubiquitous) /DB_XREF=gi:7657648 /UG=Hs.22826 tropomodulin 3 (ubiquitous) /FL=gb:AF237631.1 gb:NM_014547.1	NM_014547		NP_055362
220864_s_at	0.005852	gb:NM_015965.1 /DEF=Homo sapiens CGI-39 protein; cell death-regulatory protein GRIM19 (LOC51079), mRNA. /FEA=mRNA /GEN=LOC51079 /PROD=CGI-39 protein; cell death-regulatory protein GRIM19 /DB_XREF=gi:7705733 /UG=Hs.279574 CGI-39 protein; cell death-regulatory protein GRIM19 /FL=gb:AF132973.1 gb:AF155662.1 gb:NM_015965.1	NM_015965		NP_057049

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
220865_s_at	5.03E-04	gb:NM_014317.2 /DEF=Homo sapiens trans-prenyltransferase (TPT), mRNA. /FEA=mRNA /GEN=TPT /PROD=trans-prenyltransferase /DB_XREF=gi:11863164 /UG=Hs.279865 trans-prenyltransferase /FL=gb:NM_014317.2 gb:AF118395.1	NM_014317		NP_055132
220881_at	0.038535	gb:NM_018606.1 /DEF=Homo sapiens hypothetical protein PRO1787 (PRO1787), mRNA. /FEA=mRNA /GEN=PRO1787 /PROD=hypothetical protein PRO1787 /DB_XREF=gi:8924080 /UG=Hs.283063 hypothetical protein PRO1787 /FL=gb:AF116668.1 gb:NM_018606.1	NM_018606		NP_061076
220924_s_at	1.26E-04	gb:NM_018976.1 /DEF=Homo sapiens amino acid transporter 2 (KIAA1382), mRNA. /FEA=mRNA /GEN=KIAA1382 /PROD=amino acid transporter 2 /DB_XREF=gi:9506836 /UG=Hs.298275 amino acid transporter 2 /FL=gb:NM_018976.1	NM_018976		NP_061849
220926_s_at	3.26E-04	gb:NM_025191.1 /DEF=Homo sapiens hypothetical protein FLJ13033 (FLJ13033), mRNA. /FEA=mRNA /GEN=FLJ13033 /PROD=hypothetical protein FLJ13033 /DB_XREF=gi:13376781 /UG=Hs.301997 hypothetical protein FLJ13033 /FL=gb:NM_025191.1	NM_025191		NP_079467
220939_s_at	0.001116	gb:NM_017743.1 /DEF=Homo sapiens hypothetical protein FLJ20283 (FLJ20283), mRNA. /FEA=mRNA /GEN=FLJ20283 /PROD=hypothetical protein FLJ20283 /DB_XREF=gi:8923261 /UG=Hs.44033 hypothetical protein FLJ20283 /FL=gb:NM_017743.1	NM_017743		NP_569118

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
220940_at	2.05E-04	gb:NM_025190.1 /DEF=Homo sapiens KIAA1641 protein (KIAA1641), mRNA. /FEA=mRNA /GEN=KIAA1641 /PROD=hypothetical protein FLJ21281 /DB_XREF=gi:13449272 /UG=Hs.44566 KIAA1641 protein /FL=gb:NM_025190.1	NM_025190		NP_079466
220941_s_at	0.041055	gb:NM_017447.1 /DEF=Homo sapiens hypothetical protein LOC54149 (YG81), mRNA. /FEA=mRNA /GEN=YG81 /PROD=hypothetical protein LOC54149 /DB_XREF=gi:8394546 /UG=Hs.49391 hypothetical protein LOC54149 /FL=gb:NM_017447.1	NM_017447		NP_059143
220945_x_at	1.26E-04	gb:NM_018050.1 /DEF=Homo sapiens hypothetical protein FLJ10298 (FLJ10298), mRNA. /FEA=mRNA /GEN=FLJ10298 /PROD=hypothetical protein FLJ10298 /DB_XREF=gi:8922334 /UG=Hs.5999 hypothetical protein FLJ10298 /FL=gb:NM_018050.1	NM_018050		NP_060520
220947_s_at	7.58E-04	gb:NM_015527.1 /DEF=Homo sapiens DKFZP434P1750 protein (DKFZP434P1750), mRNA. /FEA=mRNA /GEN=DKFZP434P1750 /PROD=DKFZP434P1750 protein /DB_XREF=gi:7661587 /UG=Hs.7274 DKFZP434P1750 protein /FL=gb:NM_015527.1	NM_015527		NP_056342
220949_s_at	1.26E-04	gb:NM_024033.1 /DEF=Homo sapiens hypothetical protein MGC5242 (MGC5242), mRNA. /FEA=mRNA /GEN=MGC5242 /PROD=hypothetical protein MGC5242 /DB_XREF=gi:13162284 /UG=Hs.77365 hypothetical protein MGC5242 /FL=gb:BC000168.2 gb:NM_024033.1	NM_024033		NP_076938

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220951_s_at	0.016495	gb:NM_014576.1 /DEF=Homo sapiens Apobec-1 complementation factor; APOBEC-1 stimulating protein (ACF), mRNA. /FEA=mRNA /GEN=ACF /PROD=Apobec-1 complementation factor; APOBEC-1 stimulating protein /DB_XREF=gi:7656852 /UG=Hs.8349 Apobec-1 complementation factor; APOBEC-1 stimulating protein /FL=gb:AF209192.1 gb:NM_014576.1 gb:AF271789.1	NM_014576		NP_620311
220952_s_at	0.041055	gb:NM_019012.1 /DEF=Homo sapiens hypothetical protein (FLJ10667), mRNA. /FEA=mRNA /GEN=FLJ10667 /PROD=hypothetical protein /DB_XREF=gi:9506616 /UG=Hs.86149 phosphoinositol 3-phosphate-binding protein-2 /FL=gb:NM_019012.1	NM_019012		NP_061885
220953_s_at	0.001471	gb:NM_019061.1 /DEF=Homo sapiens hypothetical protein (FLJ20476), mRNA. /FEA=mRNA /GEN=FLJ20476 /PROD=hypothetical protein /DB_XREF=gi:9506678 /UG=Hs.93872 KIAA1682 protein /FL=gb:NM_019061.1	NM_019061		NP_061934
220956_s_at	0.026891	gb:NM_017555.1 /DEF=Homo sapiens hypothetical protein DKFZp434E026 (DKFZp434E026), mRNA. /FEA=mRNA /GEN=DKFZp434E026 /PROD=hypothetical protein DKFZp434E026 /DB_XREF=gi:8922129 /UG=Hs.9659 hypothetical protein DKFZp434E026 /FL=gb:NM_017555.1	NM_017555		NP_542770
220964_s_at	0.010205	gb:NM_030981.1 /DEF=Homo sapiens small GTP-binding protein (RAB1B), mRNA. /FEA=mRNA /GEN=RAB1B /PROD=small GTP-binding protein /DB_XREF=gi:13569961 /FL=gb:NM_030981.1	NM_030981		NP_112243

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
220980_s_at	3.26E-04	gb:NM_031284.1 /DEF=Homo sapiens hypothetical protein DKFZp434B195 (DKFZP434B195), mRNA. /FEA=mRNA /GEN=DKFZP434B195 /PROD=hypothetical protein DKFZp434B195 /DB_XREF=gi:13775193 /FL=gb:NM_031284.1	NM_031284		NP_112574
220985_s_at	0.016402	gb:NM_030954.1 /DEF=Homo sapiens hypothetical protein DKFZp564A022 (DKFZP564A022), mRNA. /FEA=mRNA /GEN=DKFZP564A022 /PROD=hypothetical protein DKFZp564A022 /DB_XREF=gi:13569925 /FL=gb:NM_030954.1	NM_030954		NP_112216
220987_s_at	0.016934	gb:NM_030952.1 /DEF=Homo sapiens hypothetical protein DKFZp434J037 (DKFZP434J037), mRNA. /FEA=mRNA /GEN=DKFZP434J037 /PROD=hypothetical protein DKFZp434J037 /DB_XREF=gi:13569921 /FL=gb:NM_030952.1	NM_030952		NP_112214
220995_at	0.00884	gb:NM_014178.1 /DEF=Homo sapiens HSPC156 protein (HSPC156), mRNA. /FEA=mRNA /GEN=HSPC156 /PROD=HSPC156 protein /DB_XREF=gi:7661811 /UG=Hs.99291 HSPC156 protein /FL=gb:AF161505.1 gb:NM_014178.1	NM_014178		NP_054897
220999_s_at	0.016934	gb:NM_030778.1 /DEF=Homo sapiens hypothetical protein PRO1331 (PRO1331), mRNA. /FEA=mRNA /GEN=PRO1331 /PROD=hypothetical protein PRO1331 /DB_XREF=gi:13562115 /FL=gb:NM_030778.1	NM_030778		NP_110405

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221002_s_at	2.05E-04	gb:NM_030927.1 /DEF=Homo sapiens hypothetical protein MGC11352 (MGC11352), mRNA. /FEA=mRNA /GEN=MGC11352 /PROD=hypothetical protein MGC11352 /DB_XREF=gi:13569888 /FL=gb:NM_030927.1	NM_030927		NP_112189
221006_s_at	7.58E-04	gb:NM_030918.1 /DEF=Homo sapiens hypothetical protein My014 (MY014), mRNA. /FEA=mRNA /GEN=MY014 /PROD=hypothetical protein My014 /DB_XREF=gi:13569876 /FL=gb:NM_030918.1	NM_030918		NP_112180
221007_s_at	0.026891	gb:NM_030917.1 /DEF=Homo sapiens hypothetical protein DKFZp586K0717 (DKFZP586K0717), mRNA. /FEA=mRNA /GEN=DKFZP586K0717 /PROD=hypothetical protein DKFZp586K0717 /DB_XREF=gi:13569873 /FL=gb:NM_030917.1	NM_030917		NP_112179
221011_s_at	3.26E-04	gb:NM_030915.1 /DEF=Homo sapiens hypothetical protein DKFZp566J091 (DKFZP566J091), mRNA. /FEA=mRNA /GEN=DKFZP566J091 /PROD=hypothetical protein DKFZp566J091 /DB_XREF=gi:13569871 /FL=gb:NM_030915.1	NM_030915		NP_112177
221012_s_at	0.016934	gb:NM_030912.1 /DEF=Homo sapiens tripartite motif protein TRIM8 (TRIM8), mRNA. /FEA=mRNA /GEN=TRIM8 /PROD=tripartite motif protein TRIM8 /DB_XREF=gi:13569865 /FL=gb:NM_030912.1	NM_030912		NP_112174

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Prot in Accession No.
221014_s_at	0.00161	gb:NM_031296.1 /DEF=Homo sapiens hypothetical protein DKFZp434G099 (DKFZP434G099), mRNA. /FEA=mRNA /GEN=DKFZP434G099 /PROD=hypothetical protein DKFZp434G099 /DB_XREF=gi:13786128 /FL=gb:NM_031296.1	NM_031296		NP_112586
221025_x_at	0.041795	gb:NM_031292.1 /DEF=Homo sapiens hypothetical protein DKFZp434G1415 (DKFZP434G1415), mRNA. /FEA=mRNA /GEN=DKFZP434G1415 /PROD=hypothetical protein DKFZp434G1415 /DB_XREF=gi:13775209 /FL=gb:NM_031292.1	NM_031292		NP_112582
221028_s_at	0.011725	gb:NM_030819.1 /DEF=Homo sapiens hypothetical protein MGC11335 (MGC11335), mRNA. /FEA=mRNA /GEN=MGC11335 /PROD=hypothetical protein MGC11335 /DB_XREF=gi:13540615 /FL=gb:NM_030819.1	NM_030819		NP_110446
221036_s_at	1.26E-04	gb:NM_031301.1 /DEF=Homo sapiens hypothetical protein DKFZp564D0372 (DKFZP564D0372), mRNA. /FEA=mRNA /GEN=DKFZP564D0372 /PROD=hypothetical protein DKFZp564D0372 /DB_XREF=gi:13775223 /FL=gb:NM_031301.1	NM_031301		NP_112591
221039_s_at	0.026891	gb:NM_018482.1 /DEF=Homo sapiens KIAA1249 protein (KIAA1249), mRNA. /FEA=mRNA /GEN=KIAA1249 /PROD=uncharacterized gastric protein ZG14P /DB_XREF=gi:8923867 /UG=Hs.10669 KIAA1249 protein /FL=gb:AF222859.1 gb:NM_018482.1	NM_018482		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221041_s_at	0.001109	gb:NM_012434.1 /DEF=Homo sapiens solute carrier family 17 (anionsugar transporter), member 5 (SLC17A5), mRNA. /FEA=mRNA /GEN=SLC17A5 /PROD=solute carrier family 17 (anionsugar transporter), member 5 /DB_XREF=gi:6912665 /UG=Hs.117865 solute carrier family 17 (anionsugar transporter), member 5 /FL=gb:NM_012434.1	NM_012434		NP_036566
221044_s_at	0.005852	gb:NM_021616.1 /DEF=Homo sapiens ring finger protein 21, interferon-responsive (RNF21), mRNA. /FEA=mRNA /GEN=RNF21 /PROD=ring finger protein 21, interferon-responsive /DB_XREF=gi:11055977 /UG=Hs.125300 ring finger protein 21, interferon-responsive /FL=gb:AB039903.1 gb:NM_021616.1	NM_021616		NP_569074
221046_s_at	0.00161	gb:NM_014170.1 /DEF=Homo sapiens HSPC135 protein (HSPC135), mRNA. /FEA=mRNA /GEN=HSPC135 /PROD=HSPC135 protein /DB_XREF=gi:7661795 /UG=Hs.127496 HSPC135 protein /FL=gb:AF161484.1 gb:NM_014170.1	NM_014170		NP_054889
221050_s_at	0.004541	gb:NM_019096.1 /DEF=Homo sapiens GTP binding protein 2 (GTPBP2), mRNA. /FEA=mRNA /GEN=GTPBP2 /PROD=GTP binding protein 2 /DB_XREF=gi:9506762 /UG=Hs.13011 GTP binding protein 2 /FL=gb:NM_019096.1	NM_019096		NP_061969

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221058_s_at	3.26E-04	gb:NM_016326.2 /DEF=Homo sapiens chemokine-like factor 3, alternatively spliced (LOC51192); mRNA. /FEA=mRNA /GEN=LOC51192 /PROD=chemokine-like factor 3, alternatively spliced /DB_XREF=gi:10092611 /UG=Hs.15159 chemokine-like factor, alternatively spliced /FL=gb:NM_016326.2 gb:AF135381.2	NM_016326		NP_058647
221059_s_at	7.58E-04	gb:NM_021615.1 /DEF=Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 6 (CHST6), mRNA. /FEA=mRNA /GEN=CHST6 /PROD=carbohydrate (N-acetylglucosamine 6-O)sulfotransferase 6 /DB_XREF=gi:11055975 /UG=Hs.157439 carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 6 /FL=gb:AF219990.1 gb:NM_021615.1	NM_021615		NP_067628
221082_s_at	0.003664	gb:NM_022477.1 /DEF=Homo sapiens hypothetical protein FLJ13556 similar to N-myc downstream regulated 3 (FLJ13556); mRNA. /FEA=mRNA /GEN=FLJ13556 /PROD=hypothetical protein FLJ13556 similar to N-mycdownstream regulated 3 /DB_XREF=gi:11968028 /UG=Hs.240615 hypothetical protein FLJ13556 similar to N-myc downstream regulated 3 /FL=gb:NM_022477.1	NM_022477		NP_114402

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221156_x_at	1.26E-04	gb:NM_004748.1 /DEF=Homo sapiens cell cycle progression 8 protein (CPR8), mRNA. /FEA=mRNA /GEN=CPR8 /PROD=cell cycle progression 8 protein /DB_XREF=gi:4758047 /UG=Hs.283753 cell cycle progression 8 protein /FL=gb:AF011794.1 gb:NM_004748.1	NM_004748		NP_004739
221188_s_at	2.05E-04	gb:NM_014430.1 /DEF=Homo sapiens cell death-inducing DFFA-like effector b (CIDEB), mRNA. /FEA=mRNA /GEN=CIDEB /PROD=cell death-inducing DFFA-like effector b /DB_XREF=gi:7656978 /UG=Hs.288835 cell death-inducing DFFA-like effector b /FL=gb:AF190901.1 gb:NM_014430.1	NM_014430		NP_055245
221190_s_at	0.013223	gb:NM_013326.1 /DEF=Homo sapiens colon cancer-associated protein Mic1 (MIC1), mRNA. /FEA=mRNA /GEN=MIC1 /PROD=colon cancer-associated protein Mic1 /DB_XREF=gi:7019454 /UG=Hs.289080 colon cancer-associated protein Mic1 /FL=gb:AF143536.1 gb:NM_013326.1	NM_013326		NP_037458
221194_s_at	0.016495	gb:NM_016125.1 /DEF=Homo sapiens PTD016 protein (LOC51136), mRNA. /FEA=mRNA /GEN=LOC51136 /PROD=PTD016 protein /DB_XREF=gi:7705842 /UG=Hs.30154 PTD016 protein /FL=gb:AF100745.1 gb:NM_016125.1	NM_016125		NP_057209
221196_x_at	0.00945	gb:NM_024332.1 /DEF=Homo sapiens c6.1A (C6.1A), mRNA. /FEA=mRNA /GEN=C6.1A /PROD=c6.1A /DB_XREF=gi:13236582 /UG=Hs.301927 c6.1A /FL=gb:BC002999.1 gb:NM_024332.1	NM_024332		NP_077308

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
221210_s_at	0.007779	gb:NM_030769.1 /DEF=Homo sapiens hypothetical protein similar to swine acylneuraminase (C1ORF13), mRNA. /FEA=mRNA /GEN=C1ORF13 /PROD=hypothetical protein similar to swine acylneuraminase /DB_XREF=gi:13540532 /FL=gb:NM_030769.1	NM_030769		NP_110396
221213_s_at	0.026884	gb:NM_017661.1 /DEF=Homo sapiens hypothetical protein FLJ20086 (FLJ20086), mRNA. /FEA=mRNA /GEN=FLJ20086 /PROD=hypothetical protein FLJ20086 /DB_XREF=gi:8923096 /UG=Hs.46821 hypothetical protein FLJ20086 /FL=gb:NM_017661.1	NM_017661		NP_060131
221214_s_at	5.03E-04	gb:NM_015537.1 /DEF=Homo sapiens DKFZP586J1624 protein (DKFZP586J1624), mRNA. /FEA=mRNA /GEN=DKFZP586J1624 /PROD=DKFZP586J1624 protein /DB_XREF=gi:7661679 /UG=Hs.4964 DKFZP586J1624 protein /FL=gb:NM_015537.1	NM_015537		NP_056352
221218_s_at	0.00228	gb:NM_022445.1 /DEF=Homo sapiens mouse thiamin pyrophosphokinase homolog (TPK1), mRNA. /FEA=mRNA /GEN=TPK1 /PROD=mouse thiamin pyrophosphokinase homolog /DB_XREF=gi:11990617 /UG=Hs.58715 thiamine pyrophosphokinase /FL=gb:NM_022445.1	NM_022445		NP_071890
221234_s_at	0.00161	gb:NM_021813.1 /DEF=Homo sapiens BTB and CNC homology 1, basic leucine zipper transcription factor 2 (BACH2), mRNA. /FEA=mRNA /GEN=BACH2 /PROD=BTB and CNC homology 1, basic leucine zipper transcription factor 2 /DB_XREF=gi:13540489 /FL=gb:NM_021813.1	NM_021813		NP_068585

Gene Identifier	p-value	Description	Gene Accession No.	Unig ne Accession No.	Protein Accession No.
221235_s_at	0.001116	gb:NM_030825.1 /DEF=Homo sapiens hypothetical protein FLJ22294 (FLJ22294), mRNA. /FEA=mRNA /GEN=FLJ22294 /PROD=hypothetical protein FLJ22294 /DB_XREF=gi:13540497 /FL=gb:NM_030825.1	NM_030825		
221249_s_at	0.003171	gb:NM_030802.1 /DEF=Homo sapiens CEBP-induced protein (LOC81558), mRNA. /FEA=mRNA /GEN=LOC81558 /PROD=CEBP-induced protein /DB_XREF=gi:13540589 /FL=gb:NM_030802.1	NM_030802		NP_110429
221257_x_at	3.26E-04	gb:NM_030793.1 /DEF=Homo sapiens hypothetical protein SP329 (SP329), mRNA. /FEA=mRNA /GEN=SP329 /PROD=hypothetical protein SP329 /DB_XREF=gi:13540573 /FL=gb:NM_030793.1	NM_030793		NP_110420
221263_s_at	7.58E-04	gb:NM_031287.1 /DEF=Homo sapiens hypothetical protein MGC3133 (MGC3133), mRNA. /FEA=mRNA /GEN=MGC3133 /PROD=hypothetical protein MGC3133 /DB_XREF=gi:13775199 /FL=gb:NM_031287.1	NM_031287		NP_112577
221264_s_at	1.26E-04	gb:NM_031214.1 /DEF=Homo sapiens hypothetical protein (AF311304), mRNA. /FEA=mRNA /GEN=AF311304 /PROD=hypothetical protein /DB_XREF=gi:13654285 /FL=gb:NM_031214.1	NM_031214		NP_112491
221269_s_at	0.041055	gb:NM_031286.1 /DEF=Homo sapiens SH3BGRL3-like protein (SH3BGRL3), mRNA. /FEA=mRNA /GEN=SH3BGRL3 /PROD=SH3BGRL3-like protein /DB_XREF=gi:13775197 /FL=gb:NM_031286.1	NM_031286		NP_112576

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
221332_at	0.041795	gb:NM_005448.1 /DEF=Homo sapiens bone morphogenetic protein 15 (BMP15), mRNA. /FEA=CDS /GEN=BMP15 /PROD=bone morphogenetic protein 15 precursor /DB_XREF=gi:4885096 /UG=Hs.247820 bone morphogenetic protein 15 /FL=gb:NM_005448.1	NM_005448		NP_005439
221381_s_at	1.26E-04	gb:NM_006792.2 /DEF=Homo sapiens mortality factor 4 (MORF4), mRNA. /FEA=CDS /GEN=MORF4 /PROD=mortality factor 4 /DB_XREF=gi:6996927 /UG=Hs.251396 mortality factor 4 /FL=gb:NM_006792.2	NM_006792		
221419_s_at	0.00228	gb:NM_013307.1 /DEF=Homo sapiens non-functional folate binding protein. (HSAF000381), mRNA. /FEA=CDS /GEN=HSAF000381 /PROD=non-functional folate binding protein /DB_XREF=gi:7019412 /FL=gb:NM_013307.1	NM_013307		
221421_s_at	0.015246	gb:NM_030955.1 /DEF=Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 12 (ADAMTS12), mRNA. /FEA=CDS /GEN=ADAMTS12 /PROD=a disintegrin-like and metalloprotease(reprolysin type) with thrombospondin type 1 motif, 12 /DB_XREF=gi:13569927 /FL=gb:NM_030955.1	NM_030955		NP_112217
221425_s_at	2.76E-04	gb:NM_030940.1 /DEF=Homo sapiens hypothetical protein MGC4276 similar to CG8198 (MGC4276), mRNA. /FEA=CDS /GEN=MGC4276 /PROD=hypothetical protein MGC4276 similar to CG8198 /DB_XREF=gi:13569910 /FL=gb:NM_030940.1	NM_030940		NP_112202

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221428_s_at	0.007779	gb:NM_030921.1 /DEF=Homo sapiens hypothetical protein DC42 (DC42), mRNA. /FEA=CDS /GEN=DC42 /PROD=hypothetical protein DC42 /DB_XREF=gi:13569880 /FL=gb:NM_030921.1	NM_030921		NP_112183
221430_s_at	0.007779	gb:NM_030963.1 /DEF=Homo sapiens hypothetical protein DKFZp434O1427 (DKFZP434O1427), mRNA. /FEA=CDS /GEN=DKFZP434O1427 /PROD=hypothetical protein DKFZp434O1427 /DB_XREF=gi:13624336 /FL=gb:NM_030963.1	NM_030963		NP_112225
221437_s_at	0.025464	gb:NM_031280.1 /DEF=Homo sapiens mitochondrial ribosomal protein S15 (MRPS15), mRNA. /FEA=CDS /GEN=MRPS15 /PROD=mitochondrial ribosomal protein S15 /DB_XREF=gi:13775191 /FL=gb:NM_031280.1	NM_031280		NP_112570
221449_s_at	5.07E-04	gb:NM_030790.1 /DEF=Homo sapiens hypothetical protein CDA08 (CDA08), mRNA. /FEA=CDS /GEN=CDA08 /PROD=hypothetical protein CDA08 /DB_XREF=gi:13540566 /FL=gb:NM_030790.1	NM_030790		NP_110417
221452_s_at	3.26E-04	gb:NM_030969.1 /DEF=Homo sapiens hypothetical protein MGC1223 (MGC1223), mRNA. /FEA=CDS /GEN=MGC1223 /PROD=hypothetical protein MGC1223 /DB_XREF=gi:13624338 /FL=gb:NM_030969.1	NM_030969		NP_112231
221471_at	1.26E-04	tumor differentially expressed 1	AW173623	Hs.272168	NP_006802
221472_at	1.26E-04	tumor differentially expressed 1	AW173623	Hs.272168	NP_006802

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
221473_x_at	0.003171	gb:U49188.1 /DEF=Human placenta (Diff33) mRNA, complete cds. /FEA=mRNA /GEN=Diff33 /DB_XREF=gi:1293562 /UG=Hs.272168 tumor differentially expressed 1 /FL=gb:U49188.1 gb:AF112227.1 gb:AF153979.1 gb:NM_006811.1	U49188		NP_006802
221474_at	0.013223	gb:U26162.1 /DEF=Human myosin regulatory light chain mRNA, complete cds. /FEA=mRNA /PROD=myosin regulatory light chain /DB_XREF=gi:829622 /UG=Hs.180224 death-associated protein 6 /FL=gb:BC004994.1 gb:D50372.1 gb:U26162.1	U26162		NP_291024
221475_s_at	0.026891	Consensus includes gb:NM_002948.1 /DEF=Homo sapiens ribosomal protein L15 (RPL15), mRNA /FEA=CDS /GEN=RPL15 /PROD=ribosomal protein L15 /DB_XREF=gi:4506602 /UG=Hs.74267 ribosomal protein L15 /FL=gb:AF279903.1 gb:L25899.1 gb:NM_002948.1	NM_002948		NP_002939
221477_s_at	1.26E-04	superoxide dismutase 2, mitochondrial	BF575213	Hs.372783	
221478_at	2.05E-04	Consensus includes gb:AL132665.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566E034 (from clone DKFZp566E034); complete cds. /FEA=mRNA /DB_XREF=gi:6137021 /UG=Hs.132955 BCL2adenovirus E1B 19kD-interacting protein 3-like /FL=gb:AF060922.1 gb:AB004788.1 gb:AF067396.1 gb:NM_004331.1 gb:AL132665.1	AL132665		NP_004322

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221479_s_at	2.05E-04	gb:AF060922.1 /DEF=Homo sapiens clone 016a05 My020 protein mRNA, complete cds. /FEA=mRNA /PROD=My020 protein /DB_XREF=gi:12001981 /UG=Hs.132955 BCL2adenovirus E1B 19kD-interacting protein 3-like /FL=gb:AF060922.1 gb:AB004788.1 gb:AF067396.1 gb:NM_004331.1 gb:AL132665.1	AF060922		NP_004322
221480_at	0.019879	heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1; 37kDa)	BG180941	Hs.303627	NP_112738
221482_s_at	0.001116	gb:BC003418.1 /DEF=Homo sapiens, cyclic AMP phosphoprotein, 19 kD, clone MGC:5468, mRNA, complete cds. /FEA=mRNA /PROD=cyclic AMP phosphoprotein, 19 kD /DB_XREF=gi:13097335 /UG=Hs.7351 cyclic AMP phosphoprotein, 19 kD /FL=gb:BC003418.1 gb:AF084555.1	BC003418		NP_006619
221484_at	0.007779	Consensus includes gb:BF691447 /FEA=EST /DB_XREF=gi:11976855 /DB_XREF=est:602247615F1 /CLONE=IMAGE:4332866 /UG=Hs.107526 UDP-Gal:betaGlcNAc: beta 1,4-galactosyltransferase, polypeptide 5 /FL=gb:AB004550.1 gb:AF038663.1 gb:NM_004776.1	NM_004776		NP_004767

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221485_at	3.26E-04	Consensus includes gb:AL035683 /DEF=Human DNA sequence from clone RP5-1063B2 on chromosome 20q13.1-13.2. Contains the 3 part of the gene for Beta-1,4-galactosyltransferase, ESTs, STSs and GSSs /FEA=mRNA /DB_XREF=gi:7288039 /UG=Hs.107526 UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 5 /FL=gb:AB004550.1 gb:AF038663.1 gb:NM_004776.1	NM_004776		NP_004767
221486_at	0.021244	gb:AF067170.1 /DEF=Homo sapiens alpha endosulfine mRNA, complete cds. /FEA=mRNA /PROD=alpha endosulfine /DB_XREF=gi:4894373 /UG=Hs.111680 endosulfine alpha /FL=gb:AF067170.1 gb:AF157510.1	AF067170		NP_004427
221489_s_at	0.021312	Consensus includes gb:W48843 /FEA=EST /DB_XREF=gi:1337022 /DB_XREF=est:zc44a09.s1 /CLONE=IMAGE:325144 /UG=Hs.285814 sprouty (Drosophila) homolog 4 /FL=gb:AF227517.1	AF227517		
221490_at	1.41E-04	gb:AL136733.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434N1010 (from clone DKFZp434N1010); complete cds. /FEA=mRNA /GEN=DKFZp434N1010 /PROD=hypothetical protein /DB_XREF=gi:12052984 /UG=Hs.75425 ubiquitin associated protein /FL=gb:AL136733.1 gb:AF222043.2 gb:NM_016525.2	AL136733		NP_057609

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Prot in Accession No.
221492_s_at	1.26E-04	gb:AF202092.1 /DEF=Homo sapiens PC3-96 mRNA, complete cds. /FEA=mRNA /PROD=PC3-96 /DB_XREF=gi:11493699 /UG=Hs.26367 PC3-96 protein /FL=gb:AF202092.1	AF202092		NP_071933
221497_x_at	0.00161	gb:BC005369.1 /DEF=Homo sapiens, chromosome 1 open reading frame 12, clone MGC:12484, mRNA, complete cds. /FEA=mRNA /PROD=chromosome 1 open reading frame 12 /DB_XREF=gi:13529208 /UG=Hs.6523 chromosome 1 open reading frame 12 /FL=gb:AF229245.1 gb:AF277176.1 gb:NM_022051.1 gb:BC005369.1	BC005369		NP_071334
221498_at	1.26E-04	ESTs, Weakly similar to hypothetical protein FLJ20378 [Homo sapiens] [H.sapiens]	BF939727	Hs.409278	NP_112180
221500_s_at	0.013223	Consensus includes gb:BE782754 /FEA=EST /DB_XREF=gi:10203952 /DB_XREF=est:601471903F1 /CLONE=IMAGE:3874921 /UG=Hs.102178 syntaxin 16 /FL=gb:AF008936.1	AF008936		NP_003754
221504_s_at	1.41E-04	gb:AF112204.1 /DEF=Homo sapiens Vacuolar proton pump subunit SFD alpha isoform mRNA, complete cds. /FEA=mRNA /PROD=Vacuolar proton pump subunit SFD alpha isoform /DB_XREF=gi:6563195 /UG=Hs.19575 CGI-11 protein /FL=gb:AF298777.1 gb:AF132945.1 gb:AF112204.1 gb:NM_015941.1	AF112204		NP_057025
221505_at	1.26E-04	leucine-rich acidic nuclear protein like	AW612574	Hs.71331	NP_112182
221506_s_at	0.020412	karyopherin beta 2b, transportin	BG258639	Hs.278378	
221507_at	6.68E-04	karyopherin beta 2b, transportin	BG258639	Hs.278378	

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
221511_x_at	1.26E-04	Consensus includes gb:AB033080.1 /DEF=Homo sapiens mRNA for KIAA1254 protein, partial cds. /FEA=mRNA /GEN=KIAA1254 /PROD=KIAA1254 protein /DB_XREF=gi:6330892 /UG=Hs.82506 KIAA1254 protein /FL=gb:AF212228.1	AB033080		NP_004739
221515_s_at	0.006295	gb:BC001214.1 /DEF=Homo sapiens, CGI-68 protein, clone MGC:682; mRNA, complete cds. /FEA=mRNA /PROD=CGI-68 protein /DB_XREF=gi:12654744 /UG=Hs.8054 CGI-68 protein /FL=gb:BC001214.1 gb:AF151826.1 gb:NM_016015.1	BC001214		NP_057393
221516_s_at	0.001116	gb:BC002587.1 /DEF=Homo sapiens, hypothetical protein; clone MGC:1067; mRNA, complete cds. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:12803520 /UG=Hs.83869 hypothetical protein /FL=gb:BC002587.1	BC002587		NP_061881
221522_at	0.041055	gb:AL136784.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L0718 (from clone DKFZp434L0718); complete cds. /FEA=mRNA /GEN=DKFZp434L0718 /PROD=hypothetical protein /DB_XREF=gi:12053080 /UG=Hs.59236 Homo sapiens mRNA; cDNA DKFZp434L0718 (from clone DKFZp434L0718); complete cds /FL=gb:AL136784.1	AL136784		NP_115515

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Acc ssion No.
221523_s_at	1.26E-04	Consensus includes gb:AL138717 /DEF=Human DNA sequence from clone RP11-11D8 on chromosome 6 Contains the 5 end of the gene for a yeast ubiquitin conjugating enzyme UBC6 homolog, the gene for a possible GTP binding protein, a NACA (nascent-polypeptide-associated complex alpha polype... /FEA=mRNA_2 /DB_XREF=gi:8894207 /UG=Hs.238679 Rag D protein /FL=gb:NM_021244.1 gb:AF272036.1 gb:BC003088.1	AL138717		
221524_s_at	4.42E-04	gb:AF272036.1 /DEF=Homo sapiens Rag D mRNA, complete cds. /FEA=mRNA /PROD=Rag D /DB_XREF=gi:11181619 /UG=Hs.238679 Rag D protein /FL=gb:NM_021244.1 gb:AF272036.1 gb:BC003088.1	AF272036		NP_067067
221528_s_at	6.42E-04	gb:BC000143.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ11656, clone MGC:5247, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ11656 /DB_XREF=gi:12652782 /UG=Hs.96560 hypothetical protein FLJ11656 /FL=gb:BC000143.1	BC000143		NP_573403

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
221539_at	7.58E-04	gb:AB044548.1 /DEF=Homo sapiens POKcl.6 mRNA, for eukaryotic translation initiation factor 4E binding protein 1, complete cds. /FEA=mRNA /GEN=POKcl.6 /PROD=eukaryotic translation initiation factor 4Ebinding protein 1 /DB_XREF=gi:11527778 /UG=Hs.71819 eukaryotic translation initiation factor 4E binding protein 1 /FL=gb:AB044548.1 gb:BC004459.1 gb:NM_004095.1 gb:L36055.1	AB044548		NP_004086
221542_s_at	0.027418	SRY (sex determining region Y)-box 5	T90773	Hs.115232	NP_009106
221543_s_at	1.26E-04	gb:AL442077.1 /DEF=Homo sapiens mRNA; cDNA DKFZp667H242 (from clone DKFZp667H242); complete cds. /FEA=mRNA /GEN=DKFZp667H242 /PROD=hypothetical protein /DB_XREF=gi:10241715 /UG=Hs.125849.chromosome 8 open reading frame 2 /FL=gb:AL442077.1 gb:NM_007175.1	AL442077		NP_009106
221547_at	1.26E-04	gb:BC000794.1 /DEF=Homo sapiens, pre-mRNA splicing factor similar to S. cerevisiae Prp18, clone MGC:5075, mRNA, complete cds. /FEA=mRNA /PROD=pre-mRNA splicing factor similar to S.cerevisiae.Prp18 /DB_XREF=gi:12653992 /UG=Hs.155244 pre-mRNA processing factor 18 /FL=gb:BC000794.1 gb:U51990.1 gb:NM_003675.1	BC000794		NP_003666

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221550_at	0.048473	gb:BC002382.1 /DEF=Homo sapiens, COX15 (yeast) homolog, cytochrome c oxidase assembly protein, clone MGC:8634, mRNA, complete cds. /FEA=mRNA /PROD=COX15 (yeast) homolog, cytochrome c oxidase assembly protein /DB_XREF=gi:12803156 /UG=Hs.226581 COX15 (yeast) homolog, cytochrome c oxidase assembly protein /FL=gb:BC002382.1 gb:AF026850.1	BC002382		NP_510870
221553_at	0.013223	gb:AL136636.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564K142 (from clone DKFZp564K142); complete cds. /FEA=mRNA /GEN=DKFZp564K142 /PROD=hypothetical protein /DB_XREF=gi:12052797 /UG=Hs.323562 Homo sapiens mRNA; cDNA DKFZp564K142 (from clone DKFZp564K142); complete cds /FL=gb:AL136636.1	AL136636		NP_115497
221558_s_at	0.016934	gb:AF288571.1 /DEF=Homo sapiens lymphoid enhancer factor-1 (LEF1) mRNA, complete cds. /FEA=mRNA /GEN=LEF1 /PROD=lymphoid enhancer factor-1 /DB_XREF=gi:9858157 /UG=Hs.44865 lymphoid enhancer binding factor-1 /FL=gb:AF198532.1 gb:NM_016269.1 gb:AF288571.1	AF288571		NP_057353

Gen Identifier	p-value	Description	Gen Acc ssion No.	Unigene Accession No.	Protein Accession No.
221561_at	0.033381	gb:L21934.2 /DEF=Homo sapiens acyl-coenzyme A: cholesterol acyltransferase mRNA, complete cds. /FEA=mRNA /PROD=acyl-coenzyme A: cholesterol acyltransferase /DB_XREF=gi:4878021 /UG=Hs.14553 sterol O-acyltransferase (acyl-Coenzyme A: cholesterol acyltransferase) 1 /FL=gb:NM_003101.1 gb:L21934.2	L21934		NP_003092
221564_at	0.010205	HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae)	AL570294	Hs.235887	NP_001526
221571_at	5.03E-04	TNF receptor-associated factor 3	AI721219	Hs.297660	NP_663778
221581_s_at	0.007779	gb:AF257135.1 /DEF=Homo sapiens WBSCR15 protein (WBSCR15) mRNA, complete cds. /FEA=mRNA /GEN=WBSCR15 /PROD=WBSCR15 protein /DB_XREF=gi:9651998 /UG=Hs.56607 Williams-Beuren syndrome chromosome region 5 /FL=gb:NM_022040.1 gb:BC001609.1 gb:AF257135.1	AF257135		NP_115853
221588_x_at	0.010205	Consensus includes gb:AI640855 /FEA=EST /DB_XREF=gi:4703964 /DB_XREF=est.wa27c11.x1 /CLONE=IMAGE:2299316 /UG=Hs.293970 methylmalonate-semialdehyde dehydrogenase /FL=gb:AF130089.1	AF130089		
221596_s_at	0.02145	gb:AL136619.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564O0523 (from clone DKFZp564O0523); complete cds. /FEA=mRNA /GEN=DKFZp564O0523 /PROD=hypothetical protein /DB_XREF=gi:12052763 /UG=Hs.21590 Homo sapiens mRNA; cDNA DKFZp564O0523 (from clone DKFZp564O0523); complete cds /FL=gb:AL136619.1	AL136619		NP_115496

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
221597_s_at	0.033381	gb:BC003080.1 /DEF=Homo sapiens, Similar to HSPC171 protein, clone MGC:770, mRNA, complete cds. /FEA=mRNA /PROD=Similar to HSPC171 protein /DB_XREF=gi:13111781 /UG=Hs.279593 HSPC171 protein /FL=gb:BC003080.1	BC003080		NP_054906
221601_s_at	0.001618	regulator of Fas-induced apoptosis	AI084226	Hs.58831	NP_005440
221613_s_at	3.26E-04	gb:AL136598.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F2116 (from clone DKFZp564F2116); complete cds. /FEA=mRNA /GEN=DKFZp564F2116 /PROD=hypothetical protein /DB_XREF=gi:13276696 /UG=Hs.83954 protein associated with PRK1 /FL=gb:AL136598.1	AL136598		NP_061879
221616_s_at	0.041055	Consensus includes gb:AF077053.1 /DEF=Homo sapiens neuronal cell death-related protein mRNA, complete cds. /FEA=mRNA /PROD=neuronal cell death-related protein /DB_XREF=gi:4689153 /UG=Hs.171723 neuronal cell death-related protein /FL=gb:AF077053.1 gb:NM_015975.1 gb:AF220509.1	AF077053		NP_057059
221620_s_at	3.26E-04	gb:AF061264.1 /DEF=Homo sapiens brain my025 mRNA, complete cds. /FEA=mRNA /PROD=brain my025 /DB_XREF=gi:12001991 /UG=Hs.40065 hypothetical protein MGC4825 /FL=gb:AF061264.1	AF061264		NP_077027

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221637_s_at	0.003171	gb:BC001434.1 /DEF=Homo sapiens, clone MGC:2477, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:2477) /DB_XREF=gi:12655158 /UG=Hs.9061 hypothetical protein MGC2477 /FL=gb:BC001434.1	BC001434		NP_077004
221647_s_at	0.02145	gb:AL136935.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586O0222 (from clone DKFZp586O0222); complete cds. /FEA=mRNA /GEN=DKFZp586O0222 /PROD=hypothetical protein /DB_XREF=gi:12053364 /UG=Hs.7393 hypothetical protein from EUROIMAGE 1987170 /FL=gb:AL136935.1	AL136935		NP_068751
221654_s_at	5.03E-04	gb:AF077040.1 /DEF=Homo sapiens SIH003 mRNA, complete cds. /FEA=mRNA /PROD=SIH003 /DB_XREF=gi:4689127 /UG=Hs.251636 ubiquitin specific protease 3 /FL=gb:AF077040.1	AF077040		NP_006528
221658_s_at	0.023856	gb:AF269133.1 /DEF=Homo sapiens novel interleukin receptor (NLR) mRNA, complete cds. /FEA=mRNA /GEN=NILR /PROD=novel interleukin receptor /DB_XREF=gi:10801190 /UG=Hs.210546 interleukin 21 receptor /FL=gb:AF269133.1	AF269133		NP_851565
221664_s_at	0.029453	gb:AF154005.1 /DEF=Homo sapiens junction adhesion molecule mRNA, complete cds. /FEA=mRNA /PROD=junction adhesion molecule /DB_XREF=gi:5457118 /UG=Hs.286218 junctional adhesion molecule /FL=gb:AF154005.1	AF154005		NP_653087

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221666_s_at	7.58E-04	gb:BC004470.1 /DEF=Homo sapiens, clone MGC:10332, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10332) /DB_XREF=gi:13325315 /UG=Hs.71869 apoptosis-associated speck-like protein containing a CARD /FL=gb:BC004470.1	BC004470		NP_660184
221669_s_at	0.026891	gb:BC001964.1 /DEF=Homo sapiens, acyl-Coenzyme A dehydrogenase family, member 8, clone MGC:4966, mRNA, complete cds. /FEA=mRNA /PROD=acyl-Coenzyme A dehydrogenase family, member 8 /DB_XREF=gi:12805020 /UG=Hs.14791 acyl-Coenzyme A dehydrogenase family, member 8 /FL=gb:BC001964.1	BC001964		NP_055199
221670_s_at	0.006966	gb:AF156888.1 /DEF=Homo sapiens LIM homeobox protein 3 isoform a (LHX3) mRNA, complete cds. /FEA=mRNA /GEN=LHX3 /PROD=LIM homeobox protein 3 isoform a /DB_XREF=gi:7110142 /UG=Hs.148427 LIM homeobox protein 3 /FL=gb:AF156888.1	AF156888		NP_835258
221675_s_at	1.26E-04	gb:AF195624.1 /DEF=Homo sapiens cholinephosphotransferase 1 beta mRNA, complete cds. /FEA=mRNA /PROD=cholinephosphotransferase 1 beta /DB_XREF=gi:9502012 /UG=Hs.171889 cholinephosphotransferase 1 /FL=gb:AF195624.1	AF195624		NP_064629
221691_x_at	0.005852	gb:AB042278.1 /DEF=Homo sapiens mRNA for nucleophosminB23.2, complete cds. /FEA=mRNA /PROD=nucleophosminB23.2 /DB_XREF=gi:13536990 /FL=gb:AB042278.1	AB042278		NP_002511

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221702_s_at	0.041055	gb:AF353992.1 /DEF=Homo sapiens BBP-like protein 2 (BLP2) mRNA, complete cds. /FEA=mRNA /GEN=BLP2 /PROD=BBP-like protein 2 /DB_XREF=gi:13625462 /FL=gb:AF353992.1	AF353992		NP_510883
221705_s_at	0.018973	gb:BC005934.1 /DEF=Homo sapiens, hypothetical protein FLJ21168, clone MGC:14549, mRNA, complete cds. /FEA=mRNA /PROD=hypothetical protein FLJ21168 /DB_XREF=gi:13543553 /FL=gb:BC005934.1	BC005934		NP_079349
221717_at	0.003638	gb:L25664.1 /DEF=Human interferon mRNA, complete cds. /FEA=CDS /PROD=interferon /DB_XREF=gi:479010 /UG=Hs.247726 Human interferon mRNA, complete cds /FL=gb:L25664.1	L25664		
221724_s_at	3.26E-04	gb:AF200738.1 /DEF=Homo sapiens C-type lectin DDB27 short form (DDB27) mRNA, complete cds. /FEA=CDS /GEN=DDB27 /PROD=C-type lectin DDB27 short form /DB_XREF=gi:11493653 /UG=Hs.115515 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 /FL=gb:AF200738.1	AF200738		NP_057268
221725_at	3.26E-04	Consensus includes gb:AI962978 /FEA=EST /DB_XREF=gi:5755691 /DB_XREF=est:wt25f05.x1 /CLONE=IMAGE:2508513 /UG=Hs.288908 WAS protein family, member 2 /FL=gb:NM_006990.1 gb:AB026542.1	NM_006990		NP_008921

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221731_x_at	0.003171	Consensus includes gb:BF218922 /FEA=EST /DB_XREF=gi:11112418 /DB_XREF=est:601885091F1 /CLONE=IMAGE:4103447 /UG=Hs.81800 chondroitin sulfate proteoglycan 2 (versican)	J02814		NP_004376
221732_at	0.029713	Consensus includes gb:AK026161.1 /DEF=Homo sapiens cDNA: FLJ22508 fis, clone HRC11772. /FEA=mRNA /DB_XREF=gi:10438922 /UG=Hs.8859 Homo sapiens, Similar to RIKEN cDNA 5830420C20 gene, clone IMAGE:3633379, mRNA, partial cds	AK026161		NP_620148
221735_at	0.046687	WD repeat endosomal protein	H04342	Hs.109778	NP_065890
221736_at	2.05E-04	KIAA1219 protein	BG236163	Hs.25431	
221738_at	0.041055	KIAA1219 protein	BG236163	Hs.25431	
221740_x_at	0.04734	Homo sapiens cDNA FLJ34306 fis, clone FEBRA2008086	BG249885	Hs.339808	NP_055649
221741_s_at	7.58E-04	match: proteins: Tr:Q9Y5A9 Tr:Q9VBZ5 Tr:O77335 Tr:Q9Y172 Tr:O04503; Human DNA sequence from clone RP5-963E22 on chromosome 20 Contains the 3' end of a novel gene similar to NY-REN-2 Antigen, 5 CpG islands, ESTs, STSS and GSSs, complete sequence.	AL096828		
221748_s_at	7.22E-04	tensin	AL046979	Hs.351432	
221749_at	7.58E-04	hypothetical protein FLJ31657	AU157915	Hs.409076	NP_689971
221752_at	1.26E-04	Consensus includes gb:AL041728 /FEA=EST /DB_XREF=gi:5421076 /DB_XREF=est:DKFZp434O0717_s1 /CLONE=DKFZp434O0717 /UG=Hs.60377 KIAA1298 protein	AB037719		NP_061857
221757_at	0.002671	Consensus includes gb:BE042976 /FEA=EST /DB_XREF=gi:8360029 /DB_XREF=est:ho31d03.x1 /CLONE=IMAGE:3038981 /UG=Hs.26670 Human PAC clone RP3-515N1 from 22q11.2-q22	AC002073		
221761_at	1.26E-04	adenylosuccinate synthase	AA628948	Hs.90011	NP_001117

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221765_at	0.001747	UDP-glucose ceramide glucosyltransferase	BF970427	Hs.395863	NP_003349
221766_s_at	0.016934	Consensus includes gb:AW246673 /FEA=EST /DB_XREF=gi:6589666 /DB_XREF=est:2821951.3prime /CLONE=IMAGE:2821951 /UG=Hs.10784 hypothetical protein FLJ20037	AL078599		
221767_x_at	2.05E-04	Consensus includes gb:AA515560 /FEA=EST /DB_XREF=gi:2255160 /DB_XREF=est:ni42c09.s1 /CLONE=IMAGE:979504 /UG=Hs.283473 hypothetical protein PRO2900	AK026138		
221774_x_at	0.041055	transcription factor (p38 interacting protein)	AW003334	Hs.396316	NP_060039
221776_s_at	0.010205	bromodomain containing 7	AI885109	Hs.279762	NP_037395
221778_at	7.58E-04	KIAA1718 protein	BE217882	Hs.222707	
221786_at	1.26E-04	Consensus includes gb:BF197222 /FEA=EST /DB_XREF=gi:11085906 /DB_XREF=est:7m88b07.x1 /CLONE=IMAGE:3561949 /UG=Hs.12342 Homo sapiens clone 24538 mRNA sequence	AF055030		NP_579866
221787_at	7.58E-04	Consensus includes gb:BF431618 /FEA=EST /DB_XREF=gi:11443732 /DB_XREF=est:7o16e03.x1 /CLONE=IMAGE:3574349 /UG=Hs.12342 Homo sapiens clone 24538 mRNA sequence	AF055030		NP_579866
221790_s_at	0.009689	LDL receptor adaptor protein	AL545035	Hs.184482	NP_056442
221798_x_at	0.003171	cadherin 1, type 1, E-cadherin (epithelial)	AI183766	Hs.356360	NP_002943
221802_s_at	0.005339	KIAA1598 protein	AU157109	Hs.23740	NP_060800
221803_s_at	1.26E-04	nuclear receptor binding factor-2	AA883074	Hs.27181	NP_110386
221819_at	0.007779	RAB35, member RAS oncogene family	BF791960	Hs.94308	NP_006852

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
221820_s_at	5.03E-04	Consensus includes gb:AK024102.1 /DEF=Homo sapiens cDNA FLJ14040 fis, clone HEMBA1005513, weakly similar to MALES-ABSENT ON THE FIRST PROTEIN (EC 2.3.1.). /FEA=mRNA /DB_XREF=gi:10436399 /UG=Hs.42343 member of MYST family histone acetyl transferases, homolog of Drosophila MOF	AK024102		NP_115564
221823_at	5.64E-04	hypothetical gene supported by AF038182; BC009203	AL565741	Hs.25925	NP_149988
221824_s_at	0.001116	hypothetical protein MGC26766	AA770170	Hs.395949	NP_659458
221829_s_at	0.001116	karyopherin (importin) beta 2	AI307759	Hs.168075	NP_694858
221830_at	1.26E-04	ESTs	AI302106	Hs.409145	NP_066361
221832_s_at	0.043705	Consensus includes gb:AV741657 /FEA=EST /DB_XREF=gi:10859238 /DB_XREF=est:AV741657 /CLONE=CBMALG01 /UG=Hs.154085 leucine zipper protein 1	AK025345		
221840_at	0.00228	ESTs	AA775177	Hs.408754	NP_569119
221853_s_at	0.004341	pM5 protein	N39536	Hs.227823	NP_775885
221856_s_at	0.003171	Consensus includes gb:AL561943 /FEA=EST /DB_XREF=gi:12909874 /DB_XREF=est:AL561943 /CLONE=CS0DB002YO04 (3 prime) /UG=Hs.3346 hypothetical protein FLJ11280	AB037811		
221864_at	0.02145	hypothetical protein MGC13024	AW517464	Hs.333488	NP_689501
221867_at	0.001911	hypothetical protein FLJ31821	BF436315	Hs.351575	NP_694574
221875_x_at	0.007779	major histocompatibility complex, class I, F	AW514210	Hs.377850	NP_061823
221892_at	0.009682	Consensus includes gb:AK024548.1 /DEF=Homo sapiens cDNA: FLJ20895 fis, clone ADKA03483. /FEA=mRNA /DB_XREF=gi:10436853 /UG=Hs.285737 Homo sapiens cDNA: FLJ20895 fis, clone ADKA03483	AK024548		
221896_s_at	0.003171	likely ortholog of mouse hypoxia induced gene 1	BE739519	Hs.7917	NP_054775
221919_at	0.033381	heterogeneous nuclear ribonucleoprotein A1	AW450929	Hs.249495	NP_112420
221932_s_at	0.005379	clone FLB4739	AA133341	Hs.294083	NP_057501

Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
221943_x_at	0.00161	ribosomal protein L38	AW303136	Hs.2017	NP_000990
221952_x_at	5.03E-04	Consensus includes gb:AB037814.1 /DEF=Homo sapiens mRNA for KIAA1393 protein, partial cds. /FEA=mRNA /GEN=KIAA1393 /PROD=KIAA1393 protein /DB_XREF=gi:7243166 /UG=Hs.33317 KIAA1393 protein	AB037814		
221954_at	0.013223	chromosome 20 open reading frame 111	AA160474	Hs.75798	NP_057554
221960_s_at	0.002354	RAB2, member RAS oncogene family	AI189609	Hs.78305	NP_002856
221962_s_at	0.03002	ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast)	AI829920	Hs.409297	NP_003335
221978_at	0.001116	major histocompatibility complex, class I, F	BE138825	Hs.377850	NP_061823
221983_at	1.26E-04	hypothetical protein MGC3035	AL040896	Hs.22412	NP_077269
221984_s_at	1.26E-04	hypothetical protein MGC3035	AL040896	Hs.22412	NP_077269
221985_at	0.010205	hypothetical protein FLJ20059	AW006750	Hs.246875	NP_060114
221992_at	0.042536	ESTs, Moderately similar to nuclear pore-complex interacting protein [Homo sapiens] [H.sapiens]	AI925734	Hs.409134	
222023_at	0.018693	Consensus includes gb:AK022014.1 /DEF=Homo sapiens cDNA FLJ11952 fis, clone HEMBB1000831, weakly similar to Homo sapiens breast cancer nuclear receptor-binding auxiliary protein (BRX) mRNA. /FEA=mRNA /DB_XREF=gi:10433327 /UG=Hs.306619 Homo sapiens cDNA FLJ11952 fis, clone HEMBB1000831, weakly similar to Homo sapiens breast cancer nuclear receptor-binding auxiliary protein (BRX) mRNA	AK022014		NP_658913

Gene Identifier	p-value	Description	Gen Acc ssion No.	Unigene Accession No.	Protein Accession No.
222029_x_at	0.025293	Consensus includes gb:NM_014260.1 /DEF=Homo sapiens HLA class II region expressed gene KE2 (HKE2), mRNA. /FEA=CDS /GEN=HKE2 /PROD=HLA class II region expressed gene KE2 /DB_XREF=gi:7657161 /UG=Hs.205736 HLA class II region expressed gene KE2 /FL=gb:NM_014260.1	NM_014260		NP_055075
222035_s_at	0.003171	poly(A) polymerase alpha	AI984479	Hs.49007	NP_116021
222036_s_at	0.0213	MCM4 minichromosome maintenance deficient 4 (S. cerevisiae)	AI859865	Hs.154443	
222039_at	0.006443	ESTs, Moderately similar to PRO0478 protein [Homo sapiens] [H.sapiens]	AA292789	Hs.375584	
222040_at	0.013088	heterogeneous nuclear ribonucleoprotein A1	AI144007	Hs.249495	NP_112420
222041_at	0.021367	ESTs	BG235929	Hs.391830	NP_001374
222052_at	0.005852	ESTs, Weakly similar to 2109260A B cell growth factor [Homo sapiens] [H.sapiens]	AA001552	Hs.59839	
222061_at	0.004908	CD58 antigen, (lymphocyte function- associated antigen 3)	AA700015	Hs.75626	
222065_s_at	0.02145	flightless I homolog (Drosophila).	AI830227	Hs.83849	NP_002009
222067_x_at	0.012818	Consensus includes gb:AL353759 /DEF=Human DNA sequence from clone RP1-221C16 on chromosome 6. Contains two genes for novel histone 4 family members, two genes for novel histone 1 family members, three genes for novel histone 2B family members, a gene for a novel histone 2A family me... /FEA=mRNA_10 /DB_XREF=gi:8745068 /UG=Hs.180779 H2B histone family, member B	AL353759		
222071_s_at	1.26E-04	ESTs, Highly similar to hypothetical protein PRO2176 [Homo sapiens] [H.sapiens]	BE552428	Hs.391502	
222073_at	0.002418	collagen, type IV, alpha 3 (Goodpasture antigen)	AI694562	Hs.530	NP_112734
222077_s_at	7.58E-04	Rac GTPase activating protein 1	AU153848	Hs.23900	NP_037409

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222082_at	0.016934	HIV-1 inducer of short transcripts binding protein; lymphoma related factor	AI568395	Hs.104640	NP_056982
222111_at	0.00161	ESTs, Moderately similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AU145293	Hs.61773	
222119_s_at	1.26E-04	Consensus includes gb:AL117620.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564K2364 (from clone DKFZp564K2364); partial cds. /FEA=mRNA /GEN=DKFZp564K2364 /PROD=hypothetical protein /DB_XREF=gi:5912200 /UG=Hs.284289 vitiligo-associated protein VIT-1	AL117620		NP_079409
222129_at	0.02145	Consensus includes gb:AK026155.1 /DEF=Homo sapiens cDNA: FLJ22502 fis, clone HRC11383. /FEA=mRNA /DB_XREF=gi:10438915 /UG=Hs.298140 Homo sapiens cDNA: FLJ22502 fis, clone HRC11383	AK026155		NP_077269
222132_s_at	0.007052	Consensus includes gb:AJ278150.1 /DEF=Homo sapiens mRNA for putative lipid kinase. /FEA=mRNA /PROD=putative lipid kinase /DB_XREF=gi:8250242 /UG=Hs.260238 hypothetical protein FLJ10842	AJ278150		NP_060708
222133_s_at	0.002418	Consensus includes gb:AK022280.1 /DEF=Homo sapiens cDNA: FLJ12218 fis, clone MAMMA1001075, moderately similar to Homo sapiens CGI-72 protein mRNA. /FEA=mRNA /DB_XREF=gi:10433640 /UG=Hs.288435 Homo sapiens cDNA: FLJ12218 fis, clone MAMMA1001075, moderately similar to Homo sapiens CGI-72 protein mRNA	AK022280		NP_057102

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222143_s_at	0.001116	Consensus includes gb:AY007098.1 /DEF=Homo sapiens clone TCCCIA00183 mRNA sequence. /FEA=mRNA /DB_XREF=gi:9955989 /UG=Hs.27556 hypothetical protein FLJ22405	AY007098		NP_071930
222148_s_at	3.26E-04	hypothetical protein FLJ11040	BF688108	Hs.14202	NP_060777
222151_s_at	0.016934	Consensus includes gb:AK023738.1 /DEF=Homo sapiens cDNA FLJ13676 fis, clone PLACE1011922, weakly similar to MYOSIN HEAVY CHAIN, NONMUSCLE TYPE B. /FEA=mRNA /DB_XREF=gi:10435760 /UG=Hs.300876 hypothetical protein FLJ13386	AK023738		NP_079456
222156_x_at	1.26E-04	Consensus includes gb:AK022459.1 /DEF=Homo sapiens cDNA FLJ12397 fis, clone MAMMA1002769, weakly similar to Homo sapiens cell cycle progression restoration 8 protein (CPR8) mRNA. /FEA=mRNA /DB_XREF=gi:10433861 /UG=Hs.82506 KIAA1254 protein	AK022459		NP_004739
222158_s_at	0.002418	Consensus includes gb:AF229834.1 /DEF=Homo sapiens apoptosis-related protein PNAS-4 (PNAS-4) mRNA, partial cds. /FEA=mRNA /GEN=PNAS-4 /PROD=apoptosis-related protein PNAS-4 /DB_XREF=gi:7229639 /UG=Hs.42409 CGI-146 protein	AF229834		NP_057160
222165_x_at	0.04734	Consensus includes gb:AK022885.1 /DEF=Homo sapiens cDNA FLJ12823 fis, clone NT2RP2002752. /FEA=mRNA /DB_XREF=gi:10434538 /UG=Hs.95867 hypothetical protein EST00098	AK022885		NP_077017

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222173_s_at	0.001253	Consensus includes gb:AK026105.1 /DEF=Homo sapiens cDNA: FLJ22452 fis, clone HRC09667. /FEA=mRNA /DB_XREF=gi:10438847 /UG=Hs.135917 hypothetical protein DKFZp761D1823	AK026105		NP_060891
222175_s_at	0.04734	Consensus includes gb:AK000003.1 /DEF=Homo sapiens mRNA for FLJ00003 protein, partial cds. /FEA=mRNA /GEN=FLJ00003 /PROD=FLJ00003 protein /DB_XREF=gi:7209306 /UG=Hs.8657 TIG-1	AK000003		NP_056973
222186_at	0.0174	Consensus includes gb:AL109684.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 27080. /FEA=mRNA /DB_XREF=gi:5689805 /UG=Hs.306329 Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 27080	AL109684		
222187_x_at	0.007934	Consensus includes gb:X78262.1 /DEF=H.sapiens mRNA for TRE5. /FEA=mRNA /DB_XREF=gi:587440 /UG=Hs.302178 H.sapiens mRNA for TRE5	X78262		
222204_s_at	0.021165	Consensus includes gb:AL110238.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566E104 (from clone DKFZp566E104); partial cds. /FEA=mRNA /GEN=DKFZp566E104 /PROD=hypothetical protein. /DB_XREF=gi:5817180 /UG=Hs.189834 DKFZP566E104 protein	AL110238		NP_060897
222207_x_at	0.010205	Consensus includes gb:AK024602.1 /DEF=Homo sapiens cDNA: FLJ20949 fis, clone ADSE01902. /FEA=mRNA /DB_XREF=gi:10436915 /UG=Hs.220255 Homo sapiens cDNA: FLJ20949 fis, clone ADSE01902	AK024602		

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222218_s_at	0.016934	Consensus includes gb:AJ400843.1 /DEF=Homo sapiens partial mRNA for immunoglobulin-like cell surface receptor FDF03-M14, soluble alternative form. /FEA=mRNA /GEN=FDF03-M14 /PROD=cell surface receptor FDF03-M14 /DB_XREF=gi:9715838 /UG=Hs.122591 paired immunoglobulin-like receptor alpha	AJ400843		NP_840057
222230_s_at	1.26E-04	Consensus includes gb:AK022248.1 /DEF=Homo sapiens cDNA FLJ12186 fis, clone MAMMA1000824, weakly similar to ACTIN. /FEA=mRNA /DB_XREF=gi:10433604 /UG=Hs.274369 uncharacterized hypothalamus protein HARP11	AK022248		
222231_s_at	1.26E-04	Consensus includes gb:AK025328.1 /DEF=Homo sapiens cDNA: FLJ21675 fis, clone COL09090, highly similar to AF119857. Homo sapiens PRO1855 mRNA. /FEA=mRNA /DB_XREF=gi:10437822 /UG=Hs.283558 hypothetical protein PRO1855	AK025328		NP_060979
222233_s_at	0.016934	Consensus includes gb:AK022922.1 /DEF=Homo sapiens cDNA: FLJ12860 fis, clone NT2RP2003559. /FEA=mRNA /DB_XREF=gi:10434591 /UG=Hs.28891 hypothetical protein FLJ11360	AK022922		NP_071932
222235_s_at	2.05E-04	match: proteins: Tr:O02330 Tr:Q9Y2J5 Tr:Q9V5I7; Human DNA sequence from clone RP1-19N1 on chromosome Xq21.33-22.3 Contains a gene for a novel protein. Contains ESTs, STSs and GSSs, complete sequence.	AL139812		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
222244_s_at	0.007779	Consensus includes gb:AK000749.1 /DEF=Homo sapiens cDNA FLJ20742 fis, clone HEP06891. /FEA=mRNA /DB_XREF=gi:7021031 /UG=Hs.52184 hypothetical protein FLJ20618	AK000749		NP_060373
222262_s_at	0.003368	Consensus includes gb:AL137750.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434E1212 (from clone DKFZp434E1212). /FEA=mRNA /DB_XREF=gi:6808386 /UG=Hs.120439 ethanolamine kinase	AL137750		NP_061108
222266_at	6.68E-04	chromosome 19 open reading frame 2	BF796940	Hs.7943	NP_604431
222294_s_at	1.26E-04	ESTs	AW971415	Hs.165337	
222311_s_at	0.04734	ESTs, Highly similar to SRA4_HUMAN CTD-binding SR-like protein RA4 [H.sapiens]	AA648521	Hs.390734	
222326_at	0.003539	ESTs	AW973834	Hs.105884	
222366_at	3.26E-04	ESTs	W86781	Hs.293736	
222368_at	1.26E-04	hypothetical protein from EUROIMAGE 1759349	AW972351	Hs.107203	NP_660274
266_s_at	2.05E-04	CD24 antigen (small cell lung carcinoma cluster 4 antigen)	NM_013230	Hs.381003	NP_037362
31799_at	0.005276	EST	AF070618	Hs.14074	
31826_at	0.026891	KIAA0674 protein	AB014574	Hs.14799	
31835_at	0.025464	histidine-rich glycoprotein precursor; Human histidine-rich glycoprotein mRNA, complete cds.	M13149		
31845_at	0.005852	E74-like factor 4 (ets domain transcription factor)	NM_001421	Hs.151139	NP_001412
31861_at	0.016934	immunoglobulin mu binding protein 2	NM_002180	Hs.1521	NP_002171
32029_at	0.012681	3-phosphoinositide dependent protein kinase-1	NM_002613	Hs.154729	NP_002604
32032_at	7.58E-04	EST, Weakly similar to T34549 probable mucin protein DKFZp434B0635.1 - human (fragment) [H.sapiens]	NM_022719	Hs.400995	NP_073210
32091_at	1.26E-04	KIAA0446 gene product	AB007915	Hs.158286	
32836_at	0.001618	1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)	U56417		NP_116130
33494_at	0.025245	electron-transferring-flavoprotein dehydrogenase	NM_004453	Hs.323468	NP_004444

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
34031_i_at	0.004341	cerebral cavernous malformations 1	NM_004912	Hs.93810	NP_004903
34206_at	0.016934	centaurin, delta 2	AB018325	Hs.21264	NP_631920
34210_at	2.05E-04	CDW52 antigen (CAMPATH-1 antigen)	N90866	Hs.276770	NP_001794
34408_at	2.05E-04	reticulon 2	NM_005619	Hs.3803	NP_005610
34858_at	0.041055	KIAA0176 protein	D79998	Hs.4935	
35254_at	7.58E-04	FLN29 gene product	NM_006700	Hs.5148	NP_006691
35776_at	0.048473	intersectin 1 (SH3 domain protein)	NM_003024	Hs.66392	NP_003015
36019_at	2.05E-04	dom-3 homolog Z (C. elegans)	L26260	Hs.444	NP_115830
36129_at	0.004341	KIAA0397 gene product	AB007857	Hs.7416	
36545_s_at	0.003171	KIAA0542 gene product	AB011114	Hs.62209	
36554_at	0.035611	Cluster Incl. Y15521:Homo sapiens ASMTL gene /cds=(0,1889) /gb=Y15521 /gi=3808147 /ug=Hs.6315 /len=1890	Y15521		
36564_at	0.016934	hypothetical protein FLJ90005	W27419	Hs.352358	
36566_at	0.028067	cystinosis, nephropathic	NM_004937	Hs.64837	NP_004928
36907_at	0.003171	mevalonate kinase (mevalonic aciduria)	NM_000431	Hs.130607	NP_000422
36920_at	0.027243	myotubular myopathy 1	NM_000252	Hs.75302	NP_000243
37012_at	0.003171	capping protein (actin filament) muscle Z-line, beta	NM_004930	Hs.333417	NP_004921
37145_at	0.033381	Homo sapiens NKG5 gene, complete cds.	M85276		
37170_at	0.003539	BMP-2 inducible kinase	NM_017593	Hs.198642	NP_060063
37232_at	0.004341	translocase of inner mitochondrial membrane 9 homolog (yeast)	NM_014749	Hs.77724	NP_055564
37278_at	0.007779	tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome)	NM_000116	Hs.79021	NP_851831
37943_at	0.00161	KIAA0321 protein	AB002319	Hs.8663	
37965_at	0.010205	parvin, beta	AA181053	Hs.8836	NP_037459
37966_at	0.005276	parvin, beta	AA187563	Hs.8836	NP_037459
37986_at	0.009179	erythropoietin receptor	NM_000121	Hs.127826	NP_000112
38241_at	0.010205	butyrophilin, subfamily 3, member A2	NM_006994	Hs.167741	NP_008925
38671_at	0.004341	KIAA0620 protein	AB014520	Hs.301685	NP_055918
38892_at	0.010205	KIAA0240 protein	D87077	Hs.196275	
38964_r_at	0.007779	Wiskott-Aldrich syndrome (eczema-thrombocytopenia)	NM_000377	Hs.2157	NP_000368
39313_at	0.016934	protein kinase, lysine deficient 1	NM_018979	Hs.184592	NP_061852
39318_at	0.018693	T-cell leukemia/lymphoma 1A	NM_021966	Hs.2484	NP_068801
396_f_at	0.020793	erythropoietin receptor	NM_000121	Hs.127826	NP_000112
39650_s_at	7.58E-04	KIAA0435 gene product	NM_014801	Hs.31438	NP_055616
39729_at	5.03E-04	Human natural killer cell enhancing factor (NKEFB) mRNA, complete cds.	NM_005809		NP_005800
39817_s_at	0.013223	putative c-Myc-responsive	NM_006443	Hs.109752	NP_006434

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
39891_at	3.26E-04	Homo sapiens cDNA FLJ37393 fis, clone BRAMY2027073, moderately similar to Fugu rubripes zinc finger protein	AI246730	Hs.126901	
40016_g_at	0.021876	KIAA0303 protein	AB002301	Hs.54985	
40020_at	0.021876	cadherin, EGF LAG seven-pass G-type receptor 3 (flamingo homolog, Drosophila)	NM_001407	Hs.55173	NP_001398
40149_at	7.58E-04	SH2-B homolog	NM_015503	Hs.15744	NP_056318
40225_at	0.026891	cyclin G associated kinase	NM_005255	Hs.153227	NP_005246
40255_at	0.007779	Cluster Incl. AC004531:Homo sapiens Chromosome 16 BAC clone CIT987SK-A-67A1 /cds=(0,1622) /gb=AC004531 /gi=3337392 /ug=Hs.155049 /len=1623	AC004531		
41047_at	0.026891	chromosome 9 open reading frame 16	AI885170	Hs.95867	NP_077017
41469_at	0.003171	elafin has been sequenced at the protein level; pre-elafin has not; its existence is assumed from its molecular weight (PAGE analysis); putative; Homo sapiens elafin precursor, gene, complete cds.	NM_002638		NP_002629
41512_at	0.026891	BRCA1 associated protein	AL042733	Hs.122764	NP_006759
41856_at	0.013223	EST	AL049370	Hs.13350	
44111_at	0.00161	vacuolar protein sorting 33B (yeast)	AI672363	Hs.26510	NP_061138
44696_at	5.03E-04	hypothetical protein FLJ10743	AA915989	Hs.3376	NP_060671
44790_s_at	0.041055	hypothetical protein FLJ21562	AI129310	Hs.288708	NP_079389
45526_g_at	3.26E-04	hypothetical protein FLJ14154	AI246641	Hs.184870	NP_079121
45828_at	5.03E-04	hypothetical protein FLJ10241	AI768100	Hs.299867	NP_060505
46270_at	2.05E-04	ubiquitin associated protein 1	AL039447	Hs.75425	NP_057609
46323_at	7.58E-04	Ca ²⁺ -dependent endoplasmic reticulum nucleoside diphosphatase	AL120741	Hs.389537	NP_620148
47069_at	0.00228	Rho GTPase activating protein 8	AA533284	Hs.102336	NP_851852
47083_at	0.02145	hypothetical protein MGC2718	AI280108	Hs.81057	NP_076972
47105_at	0.043705	hypothetical protein FLJ20399	AA886893	Hs.8575	NP_060273
47530_at	0.016934	hypothetical protein HSPC219	AA748492	Hs.9196	NP_057565
48106_at	0.010205	hypothetical protein FLJ20489	H14241	Hs.306989	NP_060312
48531_at	0.026891	TNFAIP3 interacting protein 2	AA522816	Hs.288991	NP_077285
48612_at	0.00161	hypothetical protein FLJ31821	AA225490	Hs.351575	NP_694574
48808_at	0.014715	dihydrofolate reductase	AI144299	Hs.83765	NP_000782
48825_at	0.007934	candidate tumor suppressor p33 ING1 homolog	AA887083	Hs.108183	NP_057246
49077_at	0.00228	protein phosphatase methylesterase-1	AL040538	Hs.63304	NP_057231
49306_at	0.003539	AD037 protein	AI890191	Hs.296162	NP_835281

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
49327_at	0.033381	sirtuin silent mating type information regulation 2 homolog 3 (S. cerevisiae)	AI492888	Hs.19306	NP_036371
49679_s_at	0.011862	matrix metalloproteinase 24 (membrane-inserted)	AA243774	Hs.355907	NP_006681
50314_i_at	0.005039	chromosome 20 open reading frame 27	AI761506	Hs.274422	NP_060344
51146_at	2.17E-04	hypothetical protein FLJ20477	AA203365	Hs.26994	
51176_at	0.004341	cofactor required for Sp1 transcriptional activation, subunit 8, 34kDa	AA131335	Hs.28166	NP_004260
51200_at	0.013223	hypothetical protein FLJ20850	AI744084	Hs.30783	NP_060437
51774_s_at	0.041055	ESTs, Moderately similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AW014299	Hs.237946	
52159_at	0.019761	HEMK homolog 7kb	W93807	Hs.46907	NP_057257
52169_at	0.00161	hypothetical protein FLJ90524	AI302185	Hs.388506	NP_699166
53071_s_at	0.041055	hypothetical protein FLJ22222	AI885411	Hs.54558	NP_787098
53912_at	7.58E-04	sorting nexin 11	AI668643	Hs.15827	NP_689450
55081_at	0.02145	chromosome 22 open reading frame 23	W46406	Hs.8535	NP_203744
55692_at	2.05E-04	engulfment and cell motility 2 (ced-12 homolog, C. elegans)	W22924	Hs.96560	NP_573403
55872_at	0.028893	KIAA1196 protein	AI493119	Hs.158654	
57588_at	0.046329	solute carrier family 24 (sodium/potassium/calcium exchanger), member 3	R62432	Hs.12321	
57703_at	0.035171	hypothetical protein MGC27076	N32782	Hs.13138	NP_689912
57715_at	0.026891	hypothetical protein LOC51063	W72694	Hs.241545	NP_057000
59644_at	7.58E-04	BMP-2 inducible kinase	AI735391	Hs.198642	NP_060063
59999_at	0.016934	hypoxia-inducible factor 1, alpha subunit inhibitor	W37897	Hs.14595	NP_060372
60528_at	0.033381	phospholipase A2, group IVB (cytosolic)	N71116	Hs.198161	NP_005081
60794_f_at	3.26E-04	Human zinc finger protein zfp6 (ZF6) mRNA, partial cds	AI400621	Hs.334690	
62212_at	0.042536	hypothetical protein MGC955	W37846	Hs.55608	NP_077002
632_at	0.010205	glycogen synthase kinase 3 alpha	NM_019884	Hs.118890	NP_063937
64064_at	0.007779	immune associated nucleotide 4 like 1 (mouse)	AI435089	Hs.26194	NP_060854
64371_at	0.033491	KIAA0365 gene product	AI978718	Hs.190452	
64408_s_at	3.09E-04	serologically defined breast cancer antigen NY-BR-20	AW025529	Hs.239812	
64438_at	0.00228	hypothetical protein FLJ22222	W19668	Hs.54558	NP_787098
64440_at	0.034221	interleukin 17 receptor C	AI560217	Hs.129959	NP_703193
64486_at	7.58E-04	coronin, actin-binding protein, 1B	AI341234	Hs.408995	NP_065174
64488_at	0.019879	Homo sapiens FKSG27 (FKSG27) mRNA, complete cds	AW003091	Hs.6217	
64883_at	1.26E-04	hypothetical protein MGC26706	AI744083	Hs.65406	NP_689794

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
65630_at	0.033381	Homo sapiens, Similar to RIKEN cDNA 5530601119 gene, clone MGC:9743 IMAGE:3854028, mRNA, complete cds	AI742455	Hs.28974	NP_060739
74694_s_at	0.005852	hypothetical protein FLJ23282	AA907940	Hs.170253	NP_079092
78047_s_at	0.010205	matrix metalloproteinase 24 (membrane-inserted)	AW001777	Hs.355907	NP_006681
78383_at	5.03E-04	tumor protein p53-binding protein	AI150117	Hs.179982	NP_005793
89476_r_at	0.003928	aminopeptidase-like 1	AA398062	Hs.71746	NP_078939
90610_at	0.041055	leucine-rich repeat protein, neuronal 1	AI654857	Hs.125742	NP_002310
91684_g_at	0.01185	exosome component Rrp41	AI571298	Hs.343589	NP_061910
AFFX-BioB-3_at	0.00161	J04423 E coli bioB gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	J04423		
AFFX-BioC-3_at	0.013223	J04423 E coli bioC protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423		
AFFX-BioDn-5_at	0.004341	J04423 E coli bioD gene dethiobiotin synthetase (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423		
AFFX-hum_alu_at	0.003171	***ALU WARNING: Human Alu-Sq subfamily consensus sequence.	U14573		
AFFX-HUMGAPDH/M33197_3_at	0.001116	glyceraldehyde-3-phosphate dehydrogenase	NM_002046	Hs.169476	NP_002037
AFFX-HUMRGE/M10098_3_at	0.003539	Human 18S rRNA gene, complete.	M10098		
AFFX-HUMRGE/M10098_M_at	6.68E-04	Human 18S rRNA gene, complete.	M10098		
AFFX-M27830_5_at	1.26E-04	Human 28S ribosomal RNA gene, complete cds.	M27830		
AFFX-r2-Ec-bioB-3_at	0.003171	Escherichia coli /REF=J04423 /DEF=E coli bioB gene biotin synthetase corresponding to nucleotides 2772-3004 of J04423 /LEN=1114 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	J04423		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
AFFX-r2-Ec-bioC-3_at	0.033381	Escherichia coli /REF=J04423 /DEF=E coli bioC protein corresponding to nucleotides 4609-4883 of J04423 /LEN=777 (5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423		
AFFX-r2-Ec-bioD-5_at	0.003171	Escherichia coli /REF=J04423 /DEF=E coli bioD gene: dethiobiotin synthetase corresponding to nucleotides 5024-5244 of J04423 /LEN=676 (5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423		
AFFX-r2-Hs18SrRNA-3_s_at	0.003539	Human 18S rRNA gene, complete.	M10098		
AFFX-r2-Hs18SrRNA-M_x_at	0.018693	Human 18S rRNA gene, complete.	M10098		
AFFX-r2-Hs28SrRNA-3_at	1.26E-04	Human 28S ribosomal RNA gene.	M11167		
AFFX-r2-P1-cre-3_at	0.02145	Bacteriophage /REF=X03453 /DEF=Bacteriophage P1 cre recombinase protein corresponding to nucleotides 1032-1270 of X03453 /LEN=1058 (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	X03453		

TABLE 3K					
Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
1053_at	0.013086	replication factor C (activator 1) 2, 40kDa	NM_002914	Hs.139226	NP_002905
1294_at	0.01132	ubiquitin-activating enzyme E1-like	NM_003335	Hs.16695	NP_003326
160020_at	0.03797	matrix metalloproteinase 14 (membrane-inserted)	NM_004995	Hs.2399	NP_004986
1729_at	0.002724	TNFRSF1A-associated via death domain	L41690	Hs.89862	NP_700474
179_at	0.01616	postmeiotic segregation increased 2-like 2	U38980	Hs.177548	
200006_at	0.018398	gb:NM_007262.1 /DEF=Homo sapiens RNA-binding protein regulatory subunit (DJ-1), mRNA. /FEA=mRNA /GEN=DJ-1 /PROD=RNA-binding protein regulatory subunit /DB_XREF=gi:6005748 /UG=Hs.10958 RNA-binding protein regulatory subunit /FL=gb:AF021819.1 gb:NM_007262.1 gb:D61380.1	NM_007262		NP_009193
200007_at	0.005574	gb:NM_003134.1 /DEF=Homo sapiens signal recognition particle 14kD (homologous Alu RNA-binding protein) (SRP14), mRNA. /FEA=mRNA /GEN=SRP14 /PROD=signal recognition particle 14kD (homologous AluRNA-binding protein) /DB_XREF=gi:4507210 /UG=Hs.180394 signal recognition particle 14kD (homologous Alu RNA-binding protein) /FL=gb:NM_003134.1 gb:U07857.1	NM_003134		NP_003125

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200009_at	0.011969	gb:NM_001494.2 /DEF=Homo sapiens GDP dissociation inhibitor 2 (GDI2), mRNA. /FEA=mRNA /GEN=GDI2 /PROD=GDP dissociation inhibitor 2 /DB_XREF=gi:6598322 /UG=Hs.56845 GDP dissociation inhibitor 2 /FL=gb:BC005145.1 gb:D13988.1 gb:NM_001494.2	NM_001494		NP_001485
200011_s_at	7.76E-04	gb:NM_001659.1 /DEF=Homo sapiens ADP-ribosylation factor 3 (ARF3), mRNA. /FEA=mRNA /GEN=ARF3 /PROD=ADP-ribosylation factor 3 /DB_XREF=gi:4502202 /UG=Hs.119177 ADP-ribosylation factor 3 /FL=gb:M74491.1 gb:NM_001659.1	NM_001659		NP_001650
200022_at	0.014602	gb:NM_000979.1 /DEF=Homo sapiens ribosomal protein L18 (RPL18), mRNA. /FEA=mRNA /GEN=RPL18 /PROD=ribosomal protein L18 /DB_XREF=gi:4506606 /UG=Hs.75458 ribosomal protein L18 /FL=gb:BC000374.1 gb:L11566.1 gb:NM_000979.1	NM_000979		NP_000970

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200030_s_at	0.032771	gb:NM_002635.1 /DEF=Homo sapiens solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 3 (SLC25A3), nuclear gene encoding mitochondrial protein, transcript variant 1b, mRNA. /FEA=mRNA /GEN=SLC25A3 /PROD=phosphate carrier precursor isoform 1b /DB_XREF=gi:4505774 /UG=Hs.78713 solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 3 /FL=gb:BC000998.1 gb:BC001328.1 gb:BC003504.1 gb:BC004345.1 gb:NM_002635.1	NM_002635		NP_005879
200035_at	0.007174	gb:NM_015343.1 /DEF=Homo sapiens hypothetical protein (HSA011916), mRNA. /FEA=mRNA /GEN=HSA011916 /PROD=hypothetical protein /DB_XREF=gi:7661721 /UG=Hs.84359 hypothetical protein /FL=gb:NM_015343.1	NM_015343		NP_056158
200037_s_at	0.007591	gb:NM_016587.1 /DEF=Homo sapiens heterochromatin-like protein 1 (HECH), mRNA. /FEA=mRNA /GEN=HECH /PROD=heterochromatin-like protein 1 /DB_XREF=gi:7705406 /UG=Hs.278554 heterochromatin-like protein 1 /FL=gb:AF136630.1 gb:NM_016587.1	NM_016587		NP_057671

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200039_s_at	0.003875	gb:NM_002794.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, beta type, 2 (PSMB2), mRNA. /FEA=mRNA /GEN=PSMB2 /PROD=proteasome (prosome, macropain) subunit, betatype, 2 /DB_XREF=gi:4506194. /UG=Hs.1390 proteasome (prosome, macropain) subunit, beta type, 2 /FL=gb:BC000268.1 gb:NM_002794.1 gb:D26599.1	NM_002794		NP_002785
200040_at	0.009678	gb:NM_006559.1 /DEF=Homo sapiens GAP-associated tyrosine phosphoprotein p62 (Sam68) (SAM68), mRNA. /FEA=mRNA /GEN=SAM68 /PROD=GAP-associated tyrosine phosphoprotein p62(Sam68) /DB_XREF=gi:5730026 /UG=Hs.119537 GAP-associated tyrosine phosphoprotein p62 (Sam68) /FL=gb:BC000717.1 gb:M88108.1 gb:NM_006559.1	NM_006559		NP_006550
200043_at	0.004519	gb:NM_004450.1 /DEF=Homo sapiens enhancer of rudimentary (Drosophila) homolog (ERH), mRNA. /FEA=mRNA /GEN=ERH /PROD=enhancer of rudimentary (Drosophila) homolog /DB_XREF=gi:4758301 /UG=Hs.118757 enhancer of rudimentary (Drosophila) homolog /FL=gb:D85758.1 gb:U66871.1 gb:NM_004450.1	NM_004450		NP_004441

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
200044_at	7.25E-04	gb:NM_003769.1 /DEF=Homo sapiens splicing factor, arginineserine-rich 9 (SFRS9), mRNA. /FEA=mRNA /GEN=SFRS9 /PROD=splicing factor, arginineserine-rich 9. /DB_XREF=gi:4506902 /UG=Hs.77608 splicing factor, arginineserine-rich 9 /FL=gb:U30825.1 gb:NM_003769.1	NM_003769		NP_003760
200046_at	0.004519	gb:NM_001344.1 /DEF=Homo sapiens defender against cell death 1 (DAD1), mRNA. /FEA=mRNA /GEN=DAD1 /PROD=defender against cell death 1 /DB_XREF=gi:4503252 /UG=Hs.82890 defender against cell death 1 /FL=gb:NM_001344.1 gb:D15057.1	NM_001344		NP_001335
200053_at	0.02217	gb:NM_004890.1 /DEF=Homo sapiens sperm associated antigen 7 (SPAG7), mRNA. /FEA=mRNA /GEN=SPAG7 /PROD=sperm associated antigen 7 /DB_XREF=gi:4757715 /UG=Hs.90436 sperm associated antigen 7 /FL=gb:AF047437.1 gb:NM_004890.1	NM_004890		NP_004881

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
200055_at	0.004311	gb:NM_006284.1 /DEF=Homo sapiens TATA box binding protein (TBP)-associated factor, RNA polymerase II, H, 30kD (TAF2H), mRNA. /FEA=mRNA /GEN=TAF2H /PROD=TATA box binding protein (TBP)-associated factor, RNA polymerase II, H, 30kD /DB_XREF=gi:5454105 /UG=Hs.89657 TATA box binding protein (TBP)-associated factor, RNA polymerase II, H, 30kD /FL=gb:NM_006284.1 gb:U13991.1	NM_006284		NP_006275
200056_s_at	0.018901	gb:NM_006333.1 /DEF=Homo sapiens nuclear DNA-binding protein (C1D), mRNA. /FEA=mRNA /GEN=C1D /PROD=nuclear DNA-binding protein /DB_XREF=gi:5453582 /UG=Hs.15164 nuclear DNA-binding protein /FL=gb:NM_006333.1	NM_006333		NP_775269
200059_s_at	0.003393	gb:BC001360.1 /DEF=Homo sapiens, ras homolog gene family, member A, clone MGC:2330, mRNA, complete cds. /FEA=mRNA /PROD=ras homolog gene family, member A /DB_XREF=gi:12655024 /UG=Hs.77273 ras homolog gene family, member A /FL=gb:NM_001664.1 gb:BC001360.1 gb:L25080.1	BC001360		NP_001655

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200063_s_at	9.85E-04	gb:BC002398.1 /DEF=Homo sapiens, nucleophosmin (nucleolar phosphoprotein B23, numatrin), clone MGC:8463, mRNA, complete cds. /FEA=mRNA /PROD=nucleophosmin (nucleolar phosphoprotein B23,numatrin) /DB_XREF=gi:12803184 /UG=Hs.9614 nucleophosmin (nucleolar phosphoprotein. B23, numatrin) /FL=gb:NM_002520.1 gb:BC002398.1 gb:BC003670.1 gb:M23613.1 gb:M26697.1 gb:M28699.1	BC002398		NP_002511
200067_x_at	6.66E-04	Consensus includes gb:AL078596 /DEF=Human DNA sequence from clone RP3-429G5 on chromosome 6q21-22.1. Contains the NR2E1 gene for nuclear receptor 2E1 (taillèss, TLL, TLX, XTLL), the 3' end of the SNX3 gene for sorting nexin 3, ESTs, STSs, GSSs and four predicted CpG islands /FEA=mRNA_2 /DB_XREF=gi:6010168 /UG=Hs.12102 sorting nexin 3 /FL=gb:AF062483.1 gb:AF034546.1 gb:NM_003795.1	AL078596		
200068_s_at	0.001108	gb:M94859.1 /DEF=Human calnexin mRNA, complete cds. /FEA=mRNA /PROD=calnexin /DB_XREF=gi:179831 /UG=Hs.155560 calnexin /FL=gb:NM_001746.1 gb:BC003552.1 gb:M94859.1 gb:M98452.1 gb:L10284.1 gb:L18887.1	M94859		NP_001737

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200070_at	0.003373	gb:BC001393.1 /DEF=Homo sapiens, hypothetical protein, clone MGC:782, mRNA, complete cds. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:12655084 /UG=Hs.4973 hypothetical protein /FL=gb:BC001393.1	BC001393		NP_056495
200071_at	0.009378	splicing factor 30, survival of motor neuron-related	BF224259	Hs.79968	NP_005862
200072_s_at	0.001812	gb:AF061832.1 /DEF=Homo sapiens M4 protein deletion mutant mRNA, complete cds. /FEA=mRNA /PROD=M4 protein deletion mutant /DB_XREF=gi:3126877 /UG=Hs.79024 heterogeneous nuclear ribonucleoprotein M. /FL=gb:AF061832.1	AF061832		NP_112480
200083_at	0.014263	ubiquitin specific protease 22	AA621731	Hs.12064	
200085_s_at	0.009678	Consensus includes gb:NM_007108.1 /DEF=Homo sapiens transcription elongation factor B (SIIL), polypeptide 2 (18kD, elongin B) (TCEB2), mRNA. /FEA=CDS /GEN=TCEB2 /PROD=elongin B /DB_XREF=gi:6005889 /UG=Hs.172772 transcription elongation factor B (SIIL), polypeptide 2 (18kD, elongin B) /FL=gb:NM_007108.1 gb:L42856.1	NM_007108		NP_009039
200087_s_at	0.005574	Consensus includes gb:AK024976.1 /DEF=Homo sapiens cDNA: FLJ21323 fis, clone COL02374. /FEA=mRNA /DB_XREF=gi:10437405 /UG=Hs.75914 Homo sapiens cDNA: FLJ21323 fis, clone COL02374	AK024976		NP_006806
200095_x_at	0.039831	ribosomal protein S10	AA320764	Hs.76230	NP_001005
200096_s_at	0.002692	ATPase, H+ transporting, lysosomal 9kDa, V0 subunit e	AI862255	Hs.24322	NP_003936

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
200594_x_at	0.007591	gb:NM_004501.1 /DEF=Homo sapiens heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A) (HNRPU), mRNA. /FEA=mRNA /GEN=HNRPU /PROD=heterogeneous nuclear ribonucleoprotein U(scaffold attachment factor A) /DB_XREF=gi:4758545 /UG=Hs.103804 heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A) /FL=gb:BC003367.1 gb:BC003621.1 gb:NM_004501.1	NM_004501		NP_114032
200596_s_at	0.018839	eukaryotic translation initiation factor 3, subunit 10 theta, 150/170kDa	BE614908	Hs.198899	NP_003741
200602_at	7.25E-04	gb:NM_000484.1 /DEF=Homo sapiens amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) (APP), mRNA. /FEA=mRNA /GEN=APP /PROD=amyloid beta (A4) precursor protein (proteasenexin-II, Alzheimer disease) /DB_XREF=gi:4502166 /UG=Hs.177486 amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) /FL=gb:NM_000484.1	NM_000484		NP_000475
200603_at	0.001406	Consensus includes gb:AL050038.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566J0124 (from clone DKFZp566J0124). /FEA=mRNA /DB_XREF=gi:4884279 /UG=Hs.183037 protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1) /FL=gb:M18468.1 gb:M33336.1 gb:NM_002734.1	AL050038		NP_002725

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200605_s_at	0.013803	gb:NM_002734.1 /DEF=Homo sapiens protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1) (PRKAR1A), mRNA. /FEA=mRNA /GEN=PRKAR1A /PROD=protein kinase, cAMP-dependent, regulatory, type I, alpha /DB_XREF=gi:4506062 /UG=Hs.183037 protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1) /FL=gb:M18468.1 gb:M33336.1 gb:NM_002734.1	NM_002734		NP_002725
200609_s_at	7.76E-04	gb:NM_017491.1 /DEF=Homo sapiens WD repeat domain 1 (WDR1), transcript variant 1, mRNA. /FEA=mRNA /GEN=WDR1 /PROD=WD repeat-containing protein 1, isoform 1 /DB_XREF=gi:9257256 /UG=Hs.85100 WD repeat domain 1 /FL=gb:BC000201.1 gb:BC002489.1 gb:AF020056.1 gb:AB010427.2 gb:NM_017491.1	NM_017491		NP_059830
200611_s_at	0.043867	gb:AB010427.2 /DEF=Homo sapiens mRNA for NORI-1, complete cds. /FEA=mRNA /PROD=NORI-1 /DB_XREF=gi:5103672 /UG=Hs.85100 WD repeat domain 1 /FL=gb:BC000201.1 gb:BC002489.1 gb:AF020056.1 gb:AB010427.2 gb:NM_017491.1	AB010427		NP_059830

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
200614_at	0.001406	gb:NM_004859.1 /DEF=Homo sapiens clathrin, heavy polypeptide (Hc) (CLTC), mRNA. /FEA=mRNA /GEN=CLTC /PROD=clathrin heavy chain. /DB_XREF=gi:4758011 /UG=Hs.178710 clathrin, heavy polypeptide (Hc) /FL=gb:D21260.1 gb:NM_004859.1	NM_004859		NP_004850
200615_s_at	0.026905	adaptor-related protein complex 2, beta 1 subunit	AL567295	Hs.74626	NP_001273
200618_at	6.66E-04	gb:NM_006148.1 /DEF=Homo sapiens LIM and SH3 protein 1 (LASP1), mRNA. /FEA=mRNA /GEN=LASP1 /PROD=LIM and SH3 protein 1 /DB_XREF=gi:5453709 /UG=Hs.75080 LIM and SH3 protein 1 /FL=gb:NM_006148.1	NM_006148		NP_006139
200619_at	0.001942	gb:NM_006842.1 /DEF=Homo sapiens splicing factor 3b, subunit 2, 145kD (SF3B2), mRNA. /FEA=mRNA /GEN=SF3B2. /PROD=splicing factor 3b, subunit 2, 145kD /DB_XREF=gi:5803154 /UG=Hs.75916 splicing factor 3b, subunit 2, 145kD /FL=gb:U41371.1 gb:NM_006842.1	NM_006842		
200620_at	0.005574	gb:NM_004872.1 /DEF=Homo sapiens chromosome 1 open reading frame 8 (C1ORF8), mRNA. /FEA=mRNA /GEN=C1ORF8 /PROD=chromosome 1 open reading frame 8. /DB_XREF=gi:4758571 /UG=Hs.11441 chromosome 1 open reading frame 8 /FL=gb:BC003106.1 gb:AF290615.1 gb:AF047439.1 gb:NM_004872.1	NM_004872		NP_004863

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200621_at	0.002749	gb:NM_004078.1 /DEF=Homo sapiens cysteine and glycine-rich protein 1 (CSRP1); mRNA. /FEA=mRNA /GEN=CSRP1 /PROD=cysteine and glycine-rich protein 1 /DB_XREF=gi:4758085 /UG=Hs.108080 cysteine and glycine-rich protein 1 /FL=gb:M33146.1 gb:NM_004078.1	NM_004078		NP_004069
200622_x_at	7.63E-04	calmodulin 3 (phosphorylase kinase, delta)	AV685208	Hs.334330	NP_005175
200623_s_at	0.005574	gb:NM_005184.1 /DEF=Homo sapiens calmodulin 3 (phosphorylase kinase, delta) (CALM3); mRNA. /FEA=mRNA /GEN=CALM3 /PROD=calmodulin 3 (phosphorylase kinase, delta) /DB_XREF=gi:4885108 /UG=Hs.141011 calmodulin 3 (phosphorylase kinase, delta) /FL=gb:BC005137.1 gb:J04046.1 gb:NM_005184.1	NM_005184		NP_005175
200625_s_at	0.001406	gb:NM_006367.2 /DEF=Homo sapiens adenylyl cyclase-associated protein (CAP); mRNA. /FEA=mRNA /GEN=CAP /PROD=adenylyl cyclase-associated protein /DB_XREF=gi:10938021 /UG=Hs.104125 adenylyl cyclase-associated protein /FL=gb:NM_006367.2 gb:L12168.1 gb:M98474.1	NM_006367		NP_006358

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	D scription	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200627_at	0.018248	gb:BC003005.1 /DEF=Homo sapiens, unactive progesterone receptor, 23 kD, clone MGC:4004, mRNA, complete cds. /FEA=mRNA /PROD=unactive progesterone receptor, 23 kD /DB_XREF=gi:12804292 /UG=Hs.278270 unactive progesterone receptor, 23 kD /FL=gb:BC003005.1 gb:L24804.1 gb:NM_006601.1	BC003005		NP_006592
200630_x_at	0.003976	SET translocation (myeloid leukemia-associated)	AV702810	Hs.145279	NP_003002
200632_s_at	7.63E-04	gb:NM_006096.1 /DEF=Homo sapiens N-myc downstream regulated (NDRG1), mRNA. /FEA=mRNA /GEN=NDRG1 /PROD=N-myc downstream regulated /DB_XREF=gi:5174656 /UG=Hs.75789 N-myc downstream regulated /FL=gb:BC003175.1 gb:D87953.1 gb:AF004162.1 gb:NM_006096.1	NM_006096		NP_006087
200633_at	0.024932	gb:NM_018955.1 /DEF=Homo sapiens ubiquitin B (UBB), mRNA. /FEA=mRNA /GEN=UBB /PROD=ubiquitin B /DB_XREF=gi:11024713 /UG=Hs.183842 ubiquitin B /FL=gb:NM_018955.1 gb:BC000379.1	NM_018955		NP_061828
200634_at	0.001406	gb:NM_005022.1 /DEF=Homo sapiens profilin 1 (PFN1), mRNA. /FEA=mRNA /GEN=PFN1 /PROD=profilin 1 /DB_XREF=gi:4826897 /UG=Hs.75721 profilin 1 /FL=gb:BC002475.1 gb:J03191.1 gb:NM_005022.1	NM_005022		NP_005013

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Prot in Accession No.
200639_s_at	6.66E-04	gb:NM_003406.1 /DEF=Homo sapiens tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, zeta polypeptide (YWHAZ), mRNA. /FEA=mRNA /GEN=YWHAZ /PROD=tyrosine 3-monooxygenasetryptophan5-monooxygenase activation protein, zeta polypeptide /DB_XREF=gi:4507952 /UG=Hs.75103 tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, zeta polypeptide /FL=gb:BC003623.1 gb:M86400.1 gb:NM_003406.1 gb:U28964.1	NM_003406		NP_663723
200640_at	0.001719	gb:NM_003406.1 /DEF=Homo sapiens tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, zeta polypeptide (YWHAZ), mRNA. /FEA=mRNA /GEN=YWHAZ /PROD=tyrosine 3-monooxygenasetryptophan5-monooxygenase activation protein, zeta polypeptide /DB_XREF=gi:4507952 /UG=Hs.75103 tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, zeta polypeptide /FL=gb:BC003623.1 gb:M86400.1 gb:NM_003406.1 gb:U28964.1	NM_003406		NP_663723

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
200644_at	7.25E-04	gb:NM_023009.1 /DEF=Homo sapiens macrophage myristoylated alanine-rich C kinase substrate (MACMARCKS), mRNA. /FEA=mRNA /GEN=MACMARCKS /PROD=macrophage myristoylated alanine-rich C kinase substrate /DB_XREF=gi:13491173 /UG=Hs.75061 macrophage myristoylated alanine-rich C kinase substrate /FL=gb:NM_023009.1	NM_023009		NP_075385
200650_s_at	7.25E-04	gb:NM_005566.1 /DEF=Homo sapiens lactate dehydrogenase A (LDHA), mRNA. /FEA=mRNA /GEN=LDHA /PROD=LDHA /DB_XREF=gi:5031856 /UG=Hs.2795 lactate dehydrogenase A /FL=gb:BC001829.1 gb:NM_005566.1	NM_005566		NP_005557
200653_s_at	0.023543	gb:M27319.1 /DEF=Human calmodulin mRNA, complete cds. /FEA=mRNA /PROD=calmodulin /DB_XREF=gi:179809 /UG=Hs.177656 calmodulin.1 (phosphorylase kinase, delta) /FL=gb:M27319.1 gb:NM_006888.1	M27319		NP_008819

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200657_at	0.009678	gb:NM_001152.1 /DEF=Homo sapiens solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator); member 5 (SLC25A5), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=SLC25A5 /PROD=solute carrier family 25 (mitochondrial carrier;adenine nucleotide translocator); member 5 /DB_XREF=gi:4502098 /UG=Hs.79172 solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 5 /FL=gb:J02683.1 gb:NM_001152.1	NM_001152		NP_001143
200660_at	0.003393	gb:NM_005620.1 /DEF=Homo sapiens S100 calcium-binding protein A11 (calgizzarin) (S100A11), mRNA. /FEA=mRNA /GEN=S100A11 /PROD=S100 calcium-binding protein A11 /DB_XREF=gi:5032056 /UG=Hs.256290 S100 calcium-binding protein A11 (calgizzarin) /FL=gb:D49355.1 gb:BC001410.1 gb:D50374.1 gb:NM_005620.1 gb:D38583.1	NM_005620		NP_005611
200663_at	7.63E-04	gb:NM_001780.1 /DEF=Homo sapiens CD63 antigen (melanoma 1 antigen) (CD63), mRNA. /FEA=mRNA /GEN=CD63 /PROD=CD63 antigen (melanoma 1 antigen) /DB_XREF=gi:4502678 /UG=Hs.76294 CD63 antigen (melanoma 1 antigen) /FL=gb:BC002349.1 gb:M59907.1 gb:NM_001780.1	NM_001780		NP_001771

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200667_at	0.005347	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)	BF448062	Hs.118797	NP_003331
200668_s_at	0.002603	gb:BC003395.1 /DEF=Homo sapiens, ubiquitin-conjugating enzyme E2D 3 (homologous to yeast UBC45), clone MGC:5416, mRNA, complete cds. /FEA=mRNA /PROD=ubiquitin-conjugating enzyme E2D 3 (homologous to yeast UBC45) /DB_XREF=gi:13097281 /UG=Hs.118797 ubiquitin-conjugating enzyme E2D 3 (homologous to yeast UBC45) /FL=gb:U39318.1 gb:BC003395.1 gb:NM_003340.1	BC003395		NP_003331
200669_s_at	0.002749	gb:NM_003340.1 /DEF=Homo sapiens ubiquitin-conjugating enzyme E2D 3 (homologous to yeast UBC45) (UBE2D3), mRNA. /FEA=mRNA /GEN=UBE2D3 /PROD=ubiquitin-conjugating enzyme E2D 3 (homologous to yeast UBC45) /DB_XREF=gi:4507776 /UG=Hs.118797 ubiquitin-conjugating enzyme E2D 3 (homologous to yeast UBC45) /FL=gb:U39318.1 gb:BC003395.1 gb:NM_003340.1	NM_003340		NP_003331

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200673_at	6.66E-04	gb:NM_014713.2 /DEF=Homo sapiens lysosomal-associated protein transmembrane 4 alpha (MBNT), mRNA. /FEA=mRNA /GEN=MBNT /PROD=lysosomal-associated protein transmembrane 4alpha /DB_XREF=gi:13518239 /UG=Hs.111894 lysosomal-associated protein transmembrane 4 alpha /FL=gb:BC000421.1 gb:BC003158.1 gb:NM_014713.2 gb:D14696.1	NM_014713		NP_055528
200676_s_at	0.030237	gb:NM_003347.1 /DEF=Homo sapiens ubiquitin-conjugating enzyme E2L 3 (UBE2L3), mRNA. /FEA=mRNA /GEN=UBE2L3 /PROD=ubiquitin-conjugating enzyme E2L 3 /DB_XREF=gi:4507788 /UG=Hs.108104 ubiquitin-conjugating enzyme E2L 3 /FL=gb:NM_003347.1	NM_003347		NP_003338
200677_at	6.66E-04	gb:NM_004339.2 /DEF=Homo sapiens pituitary tumor-transforming 1 interacting protein (PTTG1IP), mRNA. /FEA=mRNA /GEN=PTTG1IP /PROD=pituitary tumor-transforming protein1-interacting protein precursor /DB_XREF=gi:11038670 /UG=Hs.111126 pituitary tumor-transforming 1 interacting protein /FL=gb:NM_004339.2 gb:BC000415.1 gb:AF149785.1	NM_004339		NP_004330

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unig n Accession No.	Protein Accession No.
200680_x_at	0.00109	gb:NM_002128.1 /DEF=Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMG1), mRNA /FEA=mRNA /GEN=HMG1 /PROD=high-mobility group (nonhistone chromosomal)protein 1 /DB_XREF=gi:4504424 /UG=Hs.274472 high-mobility group (nonhistone chromosomal) protein 1 /FL=gb:BC003378.1 gb:NM_002128.1 gb:D63874.1	NM_002128		NP_002119
200682_s_at	7.63E-04	Consensus includes gb:BG531983 /FEA=EST /DB_XREF=gi:13523521 /DB_XREF=est:602561007F1 /CLONE=IMAGE:4699176 /UG=Hs.108104 ubiquitin-conjugating enzyme E2L 3 /FL=gb:NM_003347.1	NM_003347		NP_003338
200693_at	0.049392	gb:NM_006826.1 /DEF=Homo sapiens tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, theta polypeptide (YWHAQ), mRNA /FEA=mRNA /GEN=YWHAQ /PROD=tyrosine 3-monooxygenasetryptophan5-monooxygenase activation protein, theta polypeptide /DB_XREF=gi:5803226 /UG=Hs.74405 tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, theta polypeptide /FL=gb:NM_006826.1	NM_006826		NP_006817
200696_s_at	0.003842	gb:NM_000177.1 /DEF=Homo sapiens gelsolin (amyloidosis, Finnish type) (GSN), mRNA /FEA=mRNA /GEN=GSN /PROD=gelsolin (amyloidosis, Finnish type) /DB_XREF=gi:4504164 /UG=Hs.290070 gelsolin (amyloidosis, Finnish type) /FL=gb:NM_000177.1	NM_000177		NP_000168

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Prot in Accession No.
200697_at	7.25E-04	gb:NM_000188.1 /DEF=Homo sapiens hexokinase 1 (HK1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=HK1 /PROD=hexokinase 1 /DB_XREF=gi:4504390 /UG=Hs.118625 hexokinase 1 /FL=gb:M75126.1 gb:NM_000188.1	NM_000188		NP_277035
200698_at	0.001179	Consensus includes gb:AL542253 /FEA=EST /DB_XREF=gi:12874115 /DB_XREF=est:AL542253 /CLONE=CS0DE008YC08 (3 prime) /UG=Hs.118778 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2 /FL=gb:NM_006854.2	NM_006854		NP_006845
200699_at	0.024403	Consensus includes gb:BE962456 /FEA=EST /DB_XREF=gi:11765376 /DB_XREF=est:601655751R1 /CLONE=IMAGE:3846156 /UG=Hs.118778 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2 /FL=gb:NM_006854.2	NM_006854		NP_006845
200700_s_at	0.044397	gb:NM_006854.2 /DEF=Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2 (KDEL2), mRNA. /FEA=mRNA /GEN=KDEL2 /PROD=KDEL receptor 2 /DB_XREF=gi:8051609 /UG=Hs.118778 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2 /FL=gb:NM_006854.2	NM_006854		NP_006845

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
200703_at	0.023543	gb:NM_003746.1 /DEF=Homo sapiens dynein, cytoplasmic, light polypeptide (PIN), mRNA. /FEA=mRNA /GEN=PIN /PROD=dynein, cytoplasmic, light polypeptide /DB_XREF=gi:4505812 /UG=Hs.5120 dynein, cytoplasmic, light polypeptide /FL=gb:U32944.1 gb:NM_003746.1	NM_003746		NP_003737
200704_at	0.021381	gb:AB034747.1 /DEF=Homo sapiens SIMPLE mRNA for small integral membrane protein of lysosomal late endosome, complete cds. /FEA=mRNA /GEN=SIMPLE /PROD=small integral membrane protein of lysosomal late endosome /DB_XREF=gi:12862475 /UG=Hs.76507 LPS-induced TNF-alpha factor /FL=gb:AB034747.1 gb:U77396.1 gb:AF010312.1 gb:NM_004862.1	AB034747		NP_004853
200707_at	0.034333	gb:NM_002743.1 /DEF=Homo sapiens protein kinase C substrate 80K-H (PRKCSH), mRNA. /FEA=mRNA /GEN=PRKCSH /PROD=protein kinase C substrate 80K-H. /DB_XREF=gi:4506076 /UG=Hs.1432 protein kinase C substrate 80K-H /FL=gb:J03075.1 gb:NM_002743.1 gb:AF144075.1	NM_002743		NP_002734

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200709_at	7.63E-04	gb:NM_000801.1 /DEF=Homo sapiens FK506-binding protein 1A (12kD) (FKBP1A), mRNA. /FEA=mRNA /GEN=FKBP1A /PROD=FK506-binding protein 1A (12kD) /DB_XREF=gi:4503724 /UG=Hs.752 FK506-binding protein 1A (12kD) /FL=gb:BC001925.1 gb:M34539.1 gb:NM_000801.1	NM_000801		NP_463460
200712_s_at	0.01747	microtubule-associated protein, RP/EB family, member 1	AI633566	Hs.234279	NP_036457
200713_s_at	6.66E-04	gb:NM_012325.1 /DEF=Homo sapiens microtubule-associated protein, RPEB family, member 1 (MAPRE1), mRNA. /FEA=mRNA /GEN=MAPRE1 /PROD=microtubule-associated protein, RPEB family, member 1 /DB_XREF=gi:6912493 /UG=Hs.234279 microtubule-associated protein, RPEB family, member 1 /FL=gb:NM_012325.1 gb:U24166.1	NM_012325		NP_036457
200714_x_at	0.041179	gb:NM_006812.1 /DEF=Homo sapiens amplified in osteosarcoma (OS-9), mRNA. /FEA=mRNA /GEN=OS-9 /PROD=amplified in osteosarcoma /DB_XREF=gi:5803108 /UG=Hs.76228 amplified in osteosarcoma /FL=gb:U41635.1 gb:AB002806.1 gb:NM_006812.1	NM_006812		NP_006803

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
200716_x_at	0.030968	gb:NM_012423.1 /DEF=Homo sapiens ribosomal protein L13a (RPL13A), mRNA. /FEA=mRNA /GEN=RPL13A /PROD=ribosomal protein L13a /DB_XREF=gi:6912633 /UG=Hs.119122 ribosomal protein L13a /FL=gb:BC000514.1 gb:NM_012423.1	NM_012423		NP_036555
200721_s_at	0.031624	gb:NM_005736.2 /DEF=Homo sapiens ARP1 (actin-related protein 1, yeast) homolog A (centractin alpha) (ACTR1A), mRNA. /FEA=mRNA /GEN=ACTR1A /PROD=actin-related protein 1A /DB_XREF=gi:13325058 /UG=Hs.153961 ARP1 (actin-related protein 1, yeast) homolog A (centractin alpha) /FL=gb:BC000693.1 gb:NM_005736.2	NM_005736		NP_005727
200723_s_at	0.002109	gb:NM_005898.1 /DEF=Homo sapiens membrane component, chromosome 11, surface marker 1 (M11S1), mRNA. /FEA=mRNA /GEN=M11S1 /PROD=membrane component, chromosome 11, surfacemarker 1 /DB_XREF=gi:5174502 /UG=Hs.278672 membrane component, chromosome 11, surface marker 1 /FL=gb:BC001731.1 gb:NM_005898.1	NM_005898		NP_005889
200725_x_at	0.007503	gb:NM_006013.1 /DEF=Homo sapiens ribosomal protein L10 (RPL10), mRNA. /FEA=mRNA /GEN=RPL10 /PROD=ribosomal protein L10 /DB_XREF=gi:5174430 /UG=Hs.29797 ribosomal protein L10 /FL=gb:BC003358.1 gb:M73791.1 gb:M64241.1 gb:NM_006013.1	NM_006013		NP_006004

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
200728_at	0.001406	ARP2 actin-related protein 2 homolog (yeast)	AA699583	Hs.42915	NP_005713
200729_s_at	0.020369	gb:NM_005722.1 /DEF=Homo sapiens ARP2 (actin-related protein 2, yeast) homolog (ACTR2), mRNA /FEA=mRNA /GEN=ACTR2 /PROD=ARP2 (actin-related protein 2, yeast) homolog /DB_XREF=gi:5031570 /UG=Hs.42915 ARP2 (actin-related protein 2, yeast) homolog /FL=gb:AF006082.1 gb:NM_005722.1	NM_005722		NP_005713
200732_s_at	0.002603	protein tyrosine phosphatase type IVA, member 1	BF576710	Hs.227777	NP_003454
200733_s_at	0.002692	gb:U48296.1 /DEF=Homo sapiens protein tyrosine phosphatase PTPCAAX1 (hPTPCAAX1) mRNA, complete cds. /FEA=mRNA /GEN=hPTPCAAX1 /PROD=protein tyrosine phosphatase PTPCAAX1 /DB_XREF=gi:1777754 /UG=Hs.227777 protein tyrosine phosphatase type IVA, member 1 /FL=gb:U48296.1 gb:NM_003463.1	U48296		NP_003454
200734_s_at	9.85E-04	ADP-ribosylation factor 3	BG341906	Hs.119177	NP_001650
200736_s_at	0.007652	gb:NM_000581.1 /DEF=Homo sapiens glutathione peroxidase 1 (GPX1), mRNA /FEA=mRNA /GEN=GPX1 /PROD=glutathione peroxidase 1 /DB_XREF=gi:10834975 /UG=Hs.76686 glutathione peroxidase 1 /FL=gb:NM_000581.1 gb:BC000742.1 gb:M21304.1	NM_000581		NP_000572

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
200737_at	7.63E-04	gb:NM_000291.1 /DEF=Homo sapiens phosphoglycerate kinase 1 (PGK1), mRNA. /FEA=mRNA /GEN=PGK1 /PROD=phosphoglycerate kinase 1 /DB_XREF=gi:4505762 /UG=Hs.78771 phosphoglycerate kinase 1 /FL=gb:NM_000291.1	NM_000291		NP_000282
200738_s_at	6.66E-04	gb:NM_000291.1 /DEF=Homo sapiens phosphoglycerate kinase 1 (PGK1), mRNA. /FEA=mRNA /GEN=PGK1 /PROD=phosphoglycerate kinase 1 /DB_XREF=gi:4505762 /UG=Hs.78771 phosphoglycerate kinase 1 /FL=gb:NM_000291.1	NM_000291		NP_000282
200739_s_at	0.003842	SMT3 suppressor of mif two 3 homolog 1 (yeast)	BG338532	Hs.85119	NP_008867
200740_s_at	0.002724	gb:NM_006936.1 /DEF=Homo sapiens SMT3 (suppressor of mif two 3, yeast) homolog 1 (SMT3H1), mRNA. /FEA=mRNA /GEN=SMT3H1 /PROD=SMT3 (suppressor of mif two 3, yeast) homolog 1 /DB_XREF=gi:5902095 /UG=Hs.85119 SMT3 (suppressor of mif two 3, yeast) homolog 1 /FL=gb:BC000036.1 gb:NM_006936.1	NM_006936		NP_008867
200742_s_at	7.63E-04	ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)	BG231932	Hs.20478	NP_000382

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gen Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
200743_s_at	7.63E-04	gb:NM_000391.2 /DEF=Homo sapiens ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease) (CLN2), mRNA. /FEA=mRNA /GEN=CLN2 /PROD=ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease) /DB_XREF=gi:5597012 /UG=Hs.20478 ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease) /FL=gb:AF017456.1 gb:NM_000391.2	NM_000391		NP_000382
200744_s_at	0.003669	Consensus includes gb:AI741124 /FEA=EST /DB_XREF=gi:5109412 /DB_XREF=est:wg19c04.x1 /CLONE=IMAGE:2365542 /UG=Hs.215595 guanine nucleotide binding protein (G protein), beta polypeptide 1 /FL=gb:NM_002074.1 gb:BC004186.1	NM_002074		NP_002065
200746_s_at	0.00121	gb:NM_002074.1 /DEF=Homo sapiens guanine nucleotide binding protein (G protein), beta polypeptide 1 (GNB1), mRNA. /FEA=mRNA /GEN=GNB1 /PROD=guanine nucleotide binding protein (G protein), beta polypeptide 1 /DB_XREF=gi:11321584 /UG=Hs.215595 guanine nucleotide binding protein (G protein), beta polypeptide 1 /FL=gb:NM_002074.1 gb:BC004186.1	NM_002074		NP_002065

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200752_s_at	0.021616	gb:NM_005186.2 /DEF=Homo sapiens calpain 1, (mul) large subunit (CAPN1), mRNA. /FEA=mRNA /GEN=CAPN1 /PROD=calpain 1, large subunit /DB_XREF=gi:12408655 /UG=Hs.2575 calpain 1, (mul) large subunit /FL=gb:NM_005186.2	NM_005186		NP_005177
200753_x_at	0.004987	hypothetical protein ET	BE866585	Hs.73965	NP_003007
200764_s_at	0.010713	catenin (cadherin-associated protein), alpha 1, 102kDa	A1826881	Hs.178452	NP_001894
200765_x_at	0.003009	gb:NM_001903.1 /DEF=Homo sapiens catenin (cadherin-associated protein), alpha 1 (102kD) (CTNNA1), mRNA. /FEA=mRNA /GEN=CTNNA1 /PROD=catenin (cadherin-associated protein), alpha 1 (102kD) /DB_XREF=gi:4503126 /UG=Hs.178452 catenin (cadherin-associated protein), alpha 1 (102kD) /FL=gb:L23805.1 gb:NM_001903.1	NM_001903		NP_001894
200772_x_at	0.009486	prothymosin, alpha (gene sequence 28)	BF686442	Hs.250655	NP_002814
200777_s_at	0.003965	gb:NM_014670.1 /DEF=Homo sapiens KIAA0005 gene product (KIAA0005), mRNA. /FEA=mRNA /GEN=KIAA0005 /PROD=KIAA0005 gene product /DB_XREF=gi:7661849 /UG=Hs.155291 KIAA0005 gene product /FL=gb:D13630.1 gb:NM_014670.1	NM_014670		NP_055485

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200780_x_at	0.010713	gb:NM_000516.2 /DEF=Homo sapiens guanine nucleotide binding protein (G protein), alpha stimulating activity polypeptide 1 (GNAS1), mRNA. /FEA=mRNA /GEN=GNAS1 /PROD=guanine nucleotide binding protein (G protein),alpha stimulating activity polypeptide 1 /DB_XREF=gi:8659565 /UG=Hs.273385 guanine nucleotide binding protein (G protein), alpha stimulating activity polypeptide 1 /FL=gb:BC002722.1 gb:AF088185.1 gb:NM_000516.2	NM_000516		NP_536351
200781_s_at	0.018534	gb:NM_001019.1 /DEF=Homo sapiens ribosomal protein S15a (RPS15A), mRNA. /FEA=mRNA /GEN=RPS15A /PROD=ribosomal protein S15a /DB_XREF=gi:4506688 /UG=Hs.2953 ribosomal protein S15a /FL=gb:BC001697.1 gb:NM_001019.1	NM_001019		NP_001010
200782_at	6.66E-04	gb:NM_001154.2 /DEF=Homo sapiens annexin A5 (ANXA5), mRNA. /FEA=mRNA /GEN=ANXA5 /PROD=annexin V /DB_XREF=gi:4809273 /UG=Hs.300711 annexin A5 /FL=gb:BC001429.1 gb:BC004993.1 gb:M18366.1 gb:J03745.1 gb:M21731.1 gb:M19384.1 gb:D00172.1 gb:NM_001154.2	NM_001154		NP_001145

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200785_s_at	0.003842	gb:NM_002332.1 /DEF=Homo sapiens low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor) (LRP1), mRNA. /FEA=mRNA /GEN=LRP1 /PROD=low density lipoprotein-related protein 1(alpha-2-macroglobulin receptor) /DB_XREF=gi:4758685 /UG=Hs.89137 low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor) /FL=gb:NM_002332.1	NM_002332		NP_002323
200786_at	0.006722	gb:NM_002799.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, beta type, 7 (PSMB7), mRNA. /FEA=mRNA /GEN=PSMB7 /PROD=proteasome (prosome, macropain) subunit, betatype, 7 /DB_XREF=gi:4506202 /UG=Hs.118065 proteasome (prosome, macropain) subunit, beta type, 7 /FL=gb:BC000509.1 gb:D38048.1 gb:NM_002799.1	NM_002799		NP_002790
200789_at	0.006801	gb:NM_001398.1 /DEF=Homo sapiens enoyl Coenzyme A hydratase 1, peroxisomal (ECH1), mRNA. /FEA=mRNA /GEN=ECH1 /PROD=peroxisomal enoyl-coenzyme A hydratase-likeprotein /DB_XREF=gi:4503446 /UG=Hs.196176 enoyl Coenzyme A hydratase 1, peroxisomal /FL=gb:NM_001398.1 gb:U16660.1	NM_001398		NP_001389

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200791_s_at	0.001942	gb:NM_003870.1 /DEF=Homo sapiens IQ motif containing GTPase activating protein 1 (IQGAP1), mRNA. /FEA=mRNA /GEN=IQGAP1 /PROD=IQ motif containing GTPase activating protein 1 /DB_XREF=gi:4506786 /UG=Hs.1742 IQ motif containing GTPase activating protein 1 /FL=gb:NM_003870.1 gb:L33075.1	NM_003870		NP_003861
200794_x_at	0.005896	gb:NM_014764.1 /DEF=Homo sapiens DAZ associated protein 2 (DAZAP2), mRNA. /FEA=mRNA /GEN=DAZAP2 /PROD=DAZ associated protein 2 /DB_XREF=gi:7661885 /UG=Hs.75416 DAZ associated protein 2 /FL=gb:BC002334.1 gb:D31767.1 gb:NM_014764.1	NM_014764		NP_055579
200799_at	0.002603	gb:NM_005345.3 /DEF=Homo sapiens heat shock 70kD protein 1A (HSPA1A), mRNA. /FEA=mRNA /GEN=HSPA1A /PROD=heat shock 70kD protein 1A /DB_XREF=gi:5579469 /UG=Hs.8997 heat shock 70kD protein 1A /FL=gb:BC002453.1 gb:NM_005345.3	NM_005345		NP_005336
200800_s_at	0.00121	gb:NM_005345.3 /DEF=Homo sapiens heat shock 70kD protein 1A (HSPA1A), mRNA. /FEA=mRNA /GEN=HSPA1A /PROD=heat shock 70kD protein 1A /DB_XREF=gi:5579469 /UG=Hs.8997 heat shock 70kD protein 1A /FL=gb:BC002453.1 gb:NM_005345.3	NM_005345		NP_005336

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
200803_s_at	0.00158	gb:AF033095.1 /DEF=Homo sapiens testis enhanced gene transcript protein (TEGT) mRNA, complete cds. /FEA=mRNA /GEN=TEGT /PROD=testis enhanced gene transcript protein /DB_XREF=gi:2645728 /UG=Hs.74637 testis enhanced gene transcript (BAX inhibitor 1) /FL=gb:BC000916.1 gb:AF033095.1 gb:NM_003217.1	AF033095		NP_003208
200814_at	0.007693	gb:NM_006263.1 /DEF=Homo sapiens proteasome (prosome, macropain) activator subunit 1 (PA28 alpha) (PSME1), mRNA. /FEA=mRNA /GEN=PSME1 /PROD=proteasome (prosome, macropain) activator subunit 1 (PA28 alpha) /DB_XREF=gi:5453989 /UG=Hs.75348 proteasome (prosome, macropain) activator subunit 1 (PA28 alpha) /FL=gb:BC000352.1 gb:L07633.1 gb:NM_006263.1	NM_006263		NP_788955
200815_s_at	7.25E-04	gb:L13386.1 /DEF=Homo sapiens (clone 47) Miller-Dieker lissencephaly protein (LIS1) mRNA, complete cds. /FEA=mRNA /GEN=LIS1 /PROD=Miller-Dieker lissencephaly protein /DB_XREF=gi:349825 /UG=Hs.77318 platelet-activating factor acetylhydrolase, isoform Ib, alpha subunit (45kD) /FL=gb:L13385.1 gb:L13386.1 gb:NM_000430.2	L13386		NP_000421

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
200820_at	7.76E-04	gb:NM_002812.1 /DEF=Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 8 (PSMD8), mRNA. /FEA=mRNA /GEN=PSMD8 /PROD=proteasome (prosome, macropain) 26S subunit, non-ATPase, 8 /DB_XREF=gi:4506232 /UG=Hs:78466 proteasome (prosome, macropain) 26S subunit, non-ATPase, 8 /FL=gb:D38047.1 gb:BC001164.1 gb:NM_002812.1	NM_002812		NP_002803
200821_at	0.002621	gb:NM_013995.1 /DEF=Homo sapiens lysosomal-associated membrane protein 2 (LAMP2), transcript variant LAMP2B, mRNA. /FEA=mRNA /GEN=LAMP2 /PROD=lysosomal-associated membrane protein 2 precursor /DB_XREF=gi:7669502 /UG=Hs:8262 lysosomal-associated membrane protein 2 /FL=gb:U36336.1 gb:BC002965.1 gb:NM_013995.1	NM_013995		NP_054701
200822_x_at	0.003393	gb:NM_000365.1 /DEF=Homo sapiens triosephosphate isomerase 1 (TPI1), mRNA. /FEA=mRNA /GEN=TPI1 /PROD=triosephosphate isomerase 1 /DB_XREF=gi:4507644 /UG=Hs:83848 triosephosphate isomerase 1 /FL=gb:BC004230.1 gb:NM_000365.1	NM_000365		

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200823_x_at	0.008506	gb:NM_000992.1 /DEF=Homo sapiens ribosomal protein L29 (RPL29), mRNA. /FEA=mRNA /GEN=RPL29 /PROD=ribosomal protein L29 /DB_XREF=gi:4506628 /UG=Hs.183698 ribosomal protein L29 /FL=gb:U49083.1 gb:NM_000992.1 gb:U10248.1	NM_000992		NP_000983
200827_at	0.014602	gb:NM_000302.1 /DEF=Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase, Ehlers-Danlos syndrome type VI) (PLOD), mRNA. /FEA=mRNA /GEN=PLOD /PROD=procollagen-lysine 5-dioxygenase /DB_XREF=gi:4557836 /UG=Hs.75093 procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase, Ehlers-Danlos syndrome type VI) /FL=gb:L06419.1 gb:NM_000302.1	NM_000302		NP_000293
200829_x_at	0.001719	gb:NM_003457.1 /DEF=Homo sapiens zinc finger protein 207 (ZNF207), mRNA. /FEA=mRNA /GEN=ZNF207 /PROD=zinc finger protein 207 /DB_XREF=gi:4508016 /UG=Hs.62112 zinc finger protein 207 /FL=gb:AF046001.1 gb:NM_003457.1	NM_003457		NP_003448

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
200830_at	7.76E-04	gb:NM_002808.1 /DEF=Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 2 (PSMD2), mRNA. /FEA=mRNA /GEN=PSMD2 /PROD=proteasome (prosome, macropain) 26S subunit, non-ATPase, 2 /DB_XREF=gi:4506226. /UG=Hs.74619 proteasome (prosome, macropain) 26S subunit, non-ATPase, 2 /FL=gb:D78151.1 gb:BC002368.1 gb:BC002997.1 gb:NM_002808.1 gb:U18247.1 gb:U12596.1	NM_002808		NP_002799
200838_at	0.003875	gb:NM_001908.1 /DEF=Homo sapiens cathepsin B (CTSB), mRNA. /FEA=mRNA /GEN=CTSB /PROD=cathepsin B /DB_XREF=gi:4503138 /UG=Hs.297939 cathepsin B /FL=gb:M14221.1 gb:L16510.1 gb:NM_001908.1	NM_001908		NP_680093
200839_s_at	0.001406	gb:NM_001908.1 /DEF=Homo sapiens cathepsin B (CTSB), mRNA. /FEA=mRNA /GEN=CTSB /PROD=cathepsin B /DB_XREF=gi:4503138 /UG=Hs.297939 cathepsin B /FL=gb:M14221.1 gb:L16510.1 gb:NM_001908.1	NM_001908		NP_680093
200844_s_at	6.66E-04	anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phospholipase A2)	BE869583	Hs.120	NP_004896

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200845_s_at	0.014263	gb:NM_004905.1 /DEF=Homo sapiens anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phospholipase A2) (KIAA0106), mRNA. /FEA=mRNA /GEN=KIAA0106 /PROD=anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phospholipase A2) /DB_XREF=gi:4758637 /UG=Hs.120 anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phospholipase A2) /FL=gb:D14662.1 gb:NM_004905.1	NM_004905		NP_004896
200850_s_at	0.013803	gb:NM_006621.1 /DEF=Homo sapiens S-adenosylhomocysteine hydrolase-like 1 (AHCYL1), mRNA. /FEA=mRNA /GEN=AHCYL1 /PROD=S-adenosylhomocysteine hydrolase-like 1. /DB_XREF=gi:5729723 /UG=Hs.4113 S-adenosylhomocysteine hydrolase-like 1 /FL=gb:U82761.1 gb:NM_006621.1	NM_006621		NP_006612
200851_s_at	0.021227	gb:NM_014761.1 /DEF=Homo sapiens KIAA0174 gene product (KIAA0174), mRNA. /FEA=mRNA /GEN=KIAA0174 /PROD=KIAA0174 gene product /DB_XREF=gi:7661971 /UG=Hs.75824 KIAA0174 gene product /FL=gb:D79996.1 gb:BC000430.1 gb:BC004359.1 gb:NM_014761.1	NM_014761		NP_055576

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200853_at	0.001179	gb:NM_002106.1 /DEF=Homo sapiens H2A histone family, member Z (H2AFZ), mRNA. /FEA=mRNA /GEN=H2AFZ /PROD=H2A histone family, member Z /DB_XREF=gi:4504254 /UG=Hs.119192 H2A histone family, member Z /FL=gb:M37583.1 gb:NM_002106.1	NM_002106		NP_002097
200854_at	7.76E-04	Consensus includes gb:AB028970.1 /DEF=Homo sapiens mRNA for KIAA1047 protein, partial cds. /FEA=mRNA /GEN=KIAA1047 /PROD=KIAA1047 protein /DB_XREF=gi:5689430 /UG=Hs.144904 nuclear receptor co-repressor 1. /FL=gb:AF044209.1 gb:NM_006311.1	NM_006311		NP_006302
200855_at	6.66E-04	Consensus includes gb:AW771910 /FEA=EST /DB_XREF=gi:7703971 /DB_XREF=est:hn66c11.x1 /CLONE=IMAGE:3032852 /UG=Hs.144904 nuclear receptor co-repressor 1 /FL=gb:AF044209.1 gb:NM_006311.1	NM_006311		NP_006302
200857_s_at	0.006722	gb:NM_006311.1 /DEF=Homo sapiens nuclear receptor co-repressor 1 (NCOR1), mRNA. /FEA=mRNA /GEN=NCOR1 /PROD=nuclear receptor co-repressor 1 /DB_XREF=gi:5454137 /UG=Hs.144904 nuclear receptor co-repressor 1 /FL=gb:AF044209.1 gb:NM_006311.1	NM_006311		NP_006302

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200860_s_at	6.66E-04	gb:BC000779.1 /DEF=Homo sapiens, Similar to KIAA1007 protein, clone MGC:692, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA1007 protein /DB_XREF=gi:12653966 /UG=Hs.279949 KIAA1007 protein /FL=gb:BC000779.1 gb:AF110778.1 gb:NM_016284.1	BC000779		NP_057368
200861_at	0.001213	gb:NM_016284.1 /DEF=Homo sapiens KIAA1007 protein (KIAA1007), mRNA. /FEA=mRNA /GEN=KIAA1007 /PROD=KIAA1007 protein /DB_XREF=gi:7706213 /UG=Hs.279949 KIAA1007 protein /FL=gb:BC000779.1 gb:AF110778.1 gb:NM_016284.1	NM_016284		NP_057368
200863_s_at	9.85E-04	RAB11A, member RAS oncogene family	AI215102	Hs.75618	NP_004654
200864_s_at	0.002749	gb:NM_004663.1 /DEF=Homo sapiens RAB11A, member RAS oncogene family (RAB11A), mRNA. /FEA=mRNA /GEN=RAB11A /PROD=RAB11A, member RAS oncogene family /DB_XREF=gi:4758983 /UG=Hs.75618 RAB11A, member RAS oncogene family /FL=gb:AF000231.1 gb:NM_004663.1	NM_004663		NP_004654
200869_at	0.005347	gb:NM_000980.1 /DEF=Homo sapiens ribosomal protein L18a (RPL18A), mRNA. /FEA=mRNA /GEN=RPL18A /PROD=ribosomal protein L18a /DB_XREF=gi:11415025 /UG=Hs.163593 ribosomal protein L18a /FL=gb:NM_000980.1 gb:L05093.1	NM_000980		NP_000971

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200870_at	0.003616	gb:NM_007178.1 /DEF=Homo sapiens unr-interacting protein (UNRIP), mRNA. /FEA=mRNA /GEN=UNRIP /PROD=unr-interacting protein /DB_XREF=gi:6005931 /UG=Hs.3727 unr-interacting protein /FL=gb:BC000162.1 gb:AB024327.1 gb:NM_007178.1 gb:AL136691.1 gb:AF161496.1	NM_007178		NP_009109
200876_s_at	0.01063	gb:NM_002793.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, beta type, 1 (PSMB1), mRNA. /FEA=mRNA /GEN=PSMB1 /PROD=proteasome (prosome, macropain) subunit, betatype, 1 /DB_XREF=gi:4506192 /UG=Hs.75748 proteasome (prosome, macropain) subunit, beta type, 1 /FL=gb:BC000508.1 gb:NM_002793.1	NM_002793		NP_002784
200880_at	0.007693	DnaJ (Hsp40) homolog, subfamily A, member 1	AL534104	Hs.94	NP_001530
200881_s_at	6.66E-04	gb:NM_001539.1 /DEF=Homo sapiens heat shock protein, DNAJ-like 2 (HSJ2), mRNA. /FEA=mRNA /GEN=HSJ2 /PROD=heat shock protein, DNAJ-like 2 /DB_XREF=gi:4504510 /UG=Hs.94 DnaJ (Hsp40) homolog, subfamily A, member 1 /FL=gb:D13388.1 gb:L08069.1 gb:NM_001539.1	NM_001539		NP_001530

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
200886_s_at	6.66E-04	gb:NM_002629.1 /DEF=Homo sapiens phosphoglycerate mutase 1 (brain) (PGAM1), mRNA. /FEA=mRNA /GEN=PGAM1 /PROD=phosphoglycerate mutase 1 (brain) /DB_XREF=gi:4505752 /UG=Hs.181013 phosphoglycerate mutase 1 (brain) /FL=gb:BC000455.1 gb:NM_002629.1 gb:J04173.1	NM_002629		NP_002620
200889_s_at	0.001406	signal sequence receptor, alpha (translocon-associated protein alpha)	AI016620	Hs.250773	NP_003135
200891_s_at	0.03867	gb:NM_003144.2 /DEF=Homo sapiens signal sequence receptor, alpha (translocon-associated protein alpha) (SSR1), mRNA. /FEA=mRNA /GEN=SSR1 /PROD=signal sequence receptor, alpha /DB_XREF=gi:6552340 /UG=Hs.250773 signal sequence receptor, alpha (translocon-associated protein alpha) /FL=gb:AF156965.1 gb:NM_003144.2	NM_003144		NP_003135
200896_x_at	0.003875	gb:NM_004494.1 /DEF=Homo sapiens hepatoma-derived growth factor (high-mobility group protein 1-like) (HDGF), mRNA. /FEA=mRNA /GEN=HDGF /PROD=hepatoma-derived growth factor (high-mobilitygroup protein 1-like) /DB_XREF=gi:4758515 /UG=Hs.89525 hepatoma-derived growth factor (high-mobility group protein 1-like) /FL=gb:NM_004494.1 gb:D16431.1	NM_004494		NP_004485

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
200902_at	7.25E-04	gb:NM_004261.1 /DEF=Homo sapiens 15 kDa selenoprotein (SEP15), mRNA. /FEA=mRNA /GEN=SEP15 /PROD=15 kDa selenoprotein /DB_XREF=gi:4759095 /UG=Hs.90606 15 kDa selenoprotein /FL=gb:AF288991.1 gb:BC005294.1 gb:AF051894.1 gb:NM_004261.1	NM_004261		NP_004252
200909_s_at	0.013803	gb:NM_001004.1 /DEF=Homo sapiens ribosomal protein, large P2 (RPLP2), mRNA. /FEA=mRNA /GEN=RPLP2 /PROD=ribosomal protein, large P2 /DB_XREF=gi:4506670 /UG=Hs.119500 ribosomal protein, large P2 /FL=gb:BC005354.1 gb:M17887.1 gb:NM_001004.1	NM_001004		NP_000995
200914_x_at	7.63E-04	kinectin 1 (kinesin receptor)	BF589024	Hs.211577	
200918_s_at	7.63E-04	gb:NM_003139.1 /DEF=Homo sapiens signal recognition particle receptor (docking protein) (SRPR), mRNA. /FEA=mRNA /GEN=SRPR /PROD=signal recognition particle receptor (docking protein) /DB_XREF=gi:4507222 /UG=Hs.75730 signal recognition particle receptor (docking protein) /FL=gb:BC001162.1 gb:NM_003139.1	NM_003139		NP_003130
200920_s_at	7.25E-04	B-cell translocation gene 1, anti-proliferative	AL535380	Hs.77054	NP_001722

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200925_at	7.63E-04	gb:NM_004373.1 /DEF=Homo sapiens cytochrome c oxidase subunit VIa polypeptide 1 (COX6A1), mRNA. /FEA=mRNA /GEN=COX6A1 /PROD=cytochrome c oxidase subunit VIa polypeptide 1 /DB_XREF=gi:10047079 /UG=Hs.180714 cytochrome c oxidase subunit VIa polypeptide 1 /FL=gb:NM_004373.1	NM_004373		NP_004364
200927_s_at	7.63E-04	RAB14, member RAS oncogene family	AA919115	Hs.5807	NP_057406
200929_at	0.002724	gb:NM_006827.1 /DEF=Homo sapiens transmembrane trafficking protein (TMP21), mRNA. /FEA=mRNA /GEN=TMP21 /PROD=transmembrane trafficking protein /DB_XREF=gi:5803200 /UG=Hs.74137 transmembrane trafficking protein /FL=gb:BC001825.1 gb:NM_006827.1	NM_006827		NP_006818
200931_s_at	0.020369	gb:NM_014000.1 /DEF=Homo sapiens vinculin (VCL), transcript variant meta-VCL, mRNA. /FEA=mRNA /GEN=VCL /PROD=VCL isoform meta-VCL /DB_XREF=gi:7669549 /UG=Hs.75350 vinculin /FL=gb:NM_014000.1	NM_014000		NP_054706
200932_s_at	0.001004	gb:NM_006400.2 /DEF=Homo sapiens dynactin 2 (p50) (DCTN2), mRNA. /FEA=mRNA /GEN=DCTN2 /PROD=dynactin 2 /DB_XREF=gi:13259506 /UG=Hs.84153 dynactin 2 (p50) /FL=gb:U50733.1 gb:BC000718.1 gb:NM_006400.2	NM_006400		NP_006391

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200936_at	0.009329	gb:NM_000973.1 /DEF=Homo sapiens ribosomal protein L8 (RPL8), mRNA. /FEA=mRNA /GEN=RPL8 /PROD=ribosomal protein L8 /DB_XREF=gi:4506662 /UG=Hs.178551 ribosomal protein L8 /FL=gb:BC000077.1 gb:NM_000973.1	NM_000973		NP_150644
200941_at	0.014263	Consensus includes gb:AK026575.1 /DEF=Homo sapiens cDNA: FLJ22922 fis, clone KAT06722. /FEA=mRNA /DB_XREF=gi:10439459 /UG=Hs.250899 heat shock factor binding protein 1 /FL=gb:AF068754.1 gb:NM_001537.1	AK026575		NP_001528
200942_s_at	6.66E-04	gb:NM_001537.1 /DEF=Homo sapiens heat shock factor binding protein 1 (HSBP1), mRNA. /FEA=mRNA /GEN=HSBP1 /PROD=heat shock factor binding protein 1 /DB_XREF=gi:4557646 /UG=Hs.250899 heat shock factor binding protein 1 /FL=gb:AF068754.1 gb:NM_001537.1	NM_001537		NP_001528
200943_at	0.032443	gb:NM_004965.1 /DEF=Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA. /FEA=mRNA /GEN=HMG14 /PROD=high-mobility group (nonhistone chromosomal) protein 14 /DB_XREF=gi:4826757 /UG=Hs.251064 high-mobility group (nonhistone chromosomal) protein 14 /FL=gb:BC000075.1 gb:J02621.1 gb:NM_004965.1	NM_004965		NP_004956

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
200950_at	6.66E-04	gb:NM_006409.1 /DEF=Homo sapiens actin related protein 23 complex, subunit 1A (41 kD) (ARPC1A), mRNA. /FEA=mRNA /GEN=ARPC1A /PROD=actin related protein 23 complex, subunit 1A(41 kD) /DB_XREF=gi:5454077 /UG=Hs.90370 actin related protein 23 complex, subunit 1A (41 kD) /FL=gb:NM_006409.1	NM_006409		NP_006400
200953_s_at	0.044966	gb:NM_001759.1 /DEF=Homo sapiens cyclin D2 (CCND2), mRNA. /FEA=mRNA /GEN=CCND2 /PROD=cyclin D2 /DB_XREF=gi:4502616 /UG=Hs.75586 cyclin D2 /FL=gb:M90813.1 gb:D13639.1 gb:NM_001759.1	NM_001759		NP_001750
200954_at	0.041795	gb:NM_001694.1 /DEF=Homo sapiens ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 16kD (ATP6L), mRNA. /FEA=mRNA /GEN=ATP6L /PROD=ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 16kD /DB_XREF=gi:4502312 /UG=Hs.76159 ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 16kD /FL=gb:BC004537.1 gb:M62762.1 gb:NM_001694.1	NM_001694		NP_001685
200958_s_at	0.011409	gb:NM_005625.1 /DEF=Homo sapiens syndecan binding protein (syntenin) (SDCBP), mRNA. /FEA=mRNA /GEN=SDCBP /PROD=syndecan binding protein (syntenin) /DB_XREF=gi:5032082 /UG=Hs.8180 syndecan binding protein (syntenin) /FL=gb:AF000652.1 gb:NM_005625.1	NM_005625		NP_005616

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200960_x_at	0.003875	gb:NM_007096.1 /DEF=Homo sapiens clathrin, light polypeptide (Lca) (CLTA), transcript variant brain-specific, mRNA. /FEA=mRNA /GEN=CLTA /PROD=clathrin, light polypeptide A (Lca) isoform b /DB_XREF=gi:6005992 /UG=Hs.104143 clathrin, light polypeptide (Lca) /FL=gb:M20471.1 gb:NM_007096.1	NM_007096		NP_009027
200961_at	7.25E-04	gb:NM_012248.1 /DEF=Homo sapiens selenophosphate synthetase 2 (SPS2), mRNA. /FEA=mRNA /GEN=SPS2 /PROD=selenophosphate synthetase 2 /DB_XREF=gi:7657612 /UG=Hs.118725 selenophosphate synthetase 2 /FL=gb:BC002381.1 gb:U43286.1 gb:NM_012248.1	NM_012248		NP_036380
200962_at	0.039459	ribosomal protein L31	AI348010	Hs.184014	
200964_at	0.031624	gb:NM_003334.1 /DEF=Homo sapiens ubiquitin-activating enzyme E1 (A1S9T and BN75 temperature sensitivity complementing) (UBE1), mRNA. /FEA=mRNA /GEN=UBE1 /PROD=ubiquitin-activating enzyme E1 (A1S9T and BN75 temperature sensitivity complementing) /DB_XREF=gi:4507762 /UG=Hs.2055 ubiquitin-activating enzyme E1 (A1S9T and BN75 temperature sensitivity complementing) /FL=gb:M58028.1 gb:NM_003334.1	NM_003334		NP_695012

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
200968_s_at	0.007803	gb:NM_000942.1 /DEF=Homo sapiens peptidylprolyl isomerase B (cyclophilin B) (PPIB), mRNA. /FEA=mRNA /GEN=PPIB /PROD=peptidylprolyl isomerase B (cyclophilin B) /DB_XREF=gi:4758949 /UG=Hs.699 peptidylprolyl isomerase B (cyclophilin B) /FL=gb:BC001125.1 gb:M60857.1 gb:M63573.1 gb:NM_000942.1	NM_000942		NP_000933
200970_s_at	0.009378	gb:AL136807.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L1621 (from clone DKFZp434L1621); complete cds. /FEA=mRNA /GEN=DKFZp434L1621 /PROD=hypothetical protein /DB_XREF=gi:12053124 /UG=Hs.76698 stress-associated endoplasmic reticulum protein 1; ribosome associated membrane protein 4 /FL=gb:AL136807.1 gb:AF136975.1 gb:AB022427.1 gb:NM_014445.1	AL136807		NP_055260

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200971_s_at	0.002724	gb:NM_014445.1 /DEF=Homo sapiens stress-associated endoplasmic reticulum protein 1; ribosome associated membrane protein 4 (SERP1), mRNA. /FEA=mRNA /GEN=SERP1 /PROD=stress-associated endoplasmic reticulum protein1; ribosome associated membrane protein 4 /DB_XREF=gi:7657551 /UG=Hs.76698 stress-associated endoplasmic reticulum protein 1; ribosome associated membrane protein 4 /FL=gb:AL136807.1 gb:AF136975.1 gb:AB022427.1 gb:NM_014445.1	NM_014445		NP_055260
200975_at	0.002559	gb:NM_000310.1 /DEF=Homo sapiens palmitoyl-protein thioesterase 1 (ceroid-lipofuscinosis, neuronal 1, infantile) (PPT1), mRNA. /FEA=mRNA /GEN=PPT1 /PROD=palmitoyl-protein thioesterase 1(ceroid-lipofuscinosis, neuronal 1, infantile) /DB_XREF=gi:4506030 /UG=Hs.3873 palmitoyl-protein thioesterase 1 (ceroid-lipofuscinosis, neuronal 1, infantile) /FL=gb:U44772.1 gb:NM_000310.1	NM_000310		NP_000301

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Prot in Accession No.
200976_s_at	0.018901	gb:NM_006024.2 /DEF=Homo sapiens Tax1 (human T-cell leukemia virus type I) binding protein 1 (TAX1BP1), mRNA. /FEA=mRNA /GEN=TAX1BP1 /PROD=Tax1 (human T-cell leukemia virus type I)binding protein 1 /DB_XREF=gi:5803188 /UG=Hs.5437 Tax1 (human T-cell leukemia virus type I) binding protein 1 /FL=gb:U33821.2 gb:NM_006024.2 gb:AF090891.1 gb:AF268075.1	NM_006024		NP_006015
200979_at	0.006864	pyruvate dehydrogenase (lipoamide) alpha 1	BF739979	Hs.1023	
200980_s_at	0.001479	gb:NM_000284.1 /DEF=Homo sapiens pyruvate dehydrogenase (lipoamide) alpha 1 (PDHA1), mRNA. /FEA=mRNA /GEN=PDHA1 /PROD=pyruvate dehydrogenase (lipoamide) alpha 1 /DB_XREF=gi:4505684 /UG=Hs.1023 pyruvate dehydrogenase (lipoamide) alpha 1 /FL=gb:L48690.1 gb:BC002406.1 gb:J03575.1 gb:M24848.1 gb:L13318.1 gb:NM_000284.1	NM_000284		NP_000275
200981_x_at	0.003965	gb:NM_016592.1 /DEF=Homo sapiens neuroendocrine secretory protein 55 (NESP55), mRNA. /FEA=mRNA /GEN=NESP55 /PROD=neuroendocrine secretory protein 55 /DB_XREF=gi:7706588 /UG=Hs.113368 neuroendocrine secretory protein 55 /FL=gb:AF105253.1 gb:NM_016592.1	NM_016592		NP_536351

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200983_x_at	0.007652	Consensus includes gb:BF983379 /FEA=EST /DB_XREF=gi:12386191 /DB_XREF=est:602305270F1 /CLONE=IMAGE:4396576 /UG=Hs.119663 CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) /FL=gb:NM_000611.1 gb:M34671.1	NM_000611		NP_000602
200984_s_at	0.013527	Consensus includes gb:X16447.1 /DEF=Human mRNA for CD59, an LY-6-like protein regulating complement membrane attack. /FEA=mRNA /PROD=precursor polypeptide (AA -25 to 103) /DB_XREF=gi:29805 /UG=Hs.119663 CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) /FL=gb:NM_000611.1 gb:M34671.1	NM_000611		NP_000602
200985_s_at	7.63E-04	gb:NM_000611.1 /DEF=Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA. /FEA=mRNA /GEN=CD59 /PROD=CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) /DB_XREF=gi:10835164 /UG=Hs.119663 CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) /FL=gb:NM_000611.1 gb:M34671.1	NM_000611		NP_000602

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200988_s_at	0.010171	gb:NM_005789.1 /DEF=Homo sapiens proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki) (PSME3), mRNA. /FEA=mRNA /GEN=PSME3 /PROD=proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki) /DB_XREF=gi:5031996 /UG=Hs.152978 proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki) /FL=gb:NM_005789.1 gb:U11292.1	NM_005789		NP_789839
200989_at	0.013527	gb:NM_001530.1 /DEF=Homo sapiens hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) (HIF1A), mRNA. /FEA=mRNA /GEN=HIF1A /PROD=hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) /DB_XREF=gi:4504384 /UG=Hs.197540 hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) /FL=gb:U29165.1 gb:AF304431.1 gb:NM_001530.1 gb:AF207601.1 gb:AF207602.1 gb:U22431.1	NM_001530		NP_851397

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200991_s_at	0.038859	gb:NM_014748.1 /DEF=Homo sapiens KIAA0064 gene product (KIAA0064), mRNA. /FEA=mRNA /GEN=KIAA0064 /PROD=KIAA0064 gene product /DB_XREF=gi:7661889 /UG=Hs.278569 sorting nexin 17 /FL=gb:BC002524.1 gb:BC002610.1 gb:D31764.1 gb:NM_014748.1	NM_014748		NP_055563
200996_at	0.001942	gb:NM_005721.2 /DEF=Homo sapiens ARP3 (actin-related protein 3, yeast) homolog (ACTR3), mRNA. /FEA=mRNA /GEN=ACTR3 /PROD=ARP3 (actin-related protein 3, yeast) homolog /DB_XREF=gi:7262289 /UG=Hs.5321 ARP3 (actin-related protein 3; yeast) homolog /FL=gb:AF006083.1 gb:NM_005721.2	NM_005721		NP_005712
200997_at	0.01776	gb:NM_002896.1 /DEF=Homo sapiens RNA binding motif protein 4 (RBM4), mRNA. /FEA=mRNA /GEN=RBM4 /PROD=RNA binding motif protein 4 /DB_XREF=gi:4506444 /UG=Hs.6106 RNA binding motif protein 4 /FL=gb:BC000307.1 gb:U89505.1 gb:NM_002896.1	NM_002896		NP_002887

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
200999_s_at	0.002559	gb:NM_006825.1 /DEF=Homo sapiens transmembrane protein (63kD), endoplasmic reticulumGolgi intermediate compartment (P63), mRNA. /FEA=mRNA /GEN=P63 /PROD=transmembrane protein (63kD), endoplasmicreticulumGolgi intermediate compartment /DB_XREF=gi:5803112 /UG=Hs.74368 transmembrane protein (63kD), endoplasmic reticulumGolgi intermediate compartment /FL=gb:NM_006825.1	NM_006825		NP_006816
201002_s_at	0.027219	gb:U39361.1 /DEF=Homo sapiens DNA-binding protein (CROC-1B) mRNA, complete cds. /FEA=mRNA /GEN=CROC-1B /PROD=DNA-binding protein /DB_XREF=gi:1066081 /UG=Hs.75875 ubiquitin-conjugating enzyme E2 variant 1 /FL=gb:U39361.1 gb:NM_003349.2 gb:BC000468.1	U39361		NP_071887

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201007_at	0.001812	gb:NM_000183.1 /DEF=Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase3-ketoacyl-Coenzyme A thiolaseenoyl-Coenzyme A hydratase (trifunctional protein), beta subunit (HADHB), mRNA. /FEA=mRNA /GEN=HADHB /PROD=hydroxyacyl-Coenzyme A dehydrogenase3-ketoacyl-Coenzyme A thiolaseenoyl-Coenzyme A hydratase (trifunctional protein), beta subunit /DB_XREF=gi:4504326 /UG=Hs.146812 hydroxyacyl-Coenzyme A dehydrogenase3-ketoacyl-Coenzyme A thiolaseenoyl-Coenzyme A hydratase (trifunctional protein), beta subunit /FL=gb:AF113209.1 gb:NM_000183.1 gb:D16481.1	NM_000183		NP_000174
201009_s_at	0.007652	Consensus includes gb:AI439556 /FEA=EST /DB_XREF=gi:4305149 /DB_XREF=est:tc90c12.x1 /CLONE=IMAGE:2073430 /UG=Hs.179526 upregulated by 1,25-dihydroxyvitamin D-3 /FL=gb:NM_006472.1 gb:S73591.1	NM_006472		NP_006463
201011_at	0.013021	gb:NM_002950.1 /DEF=Homo sapiens ribophorin I (RPN1), mRNA. /FEA=mRNA /GEN=RPN1 /PROD=ribophorin I /DB_XREF=gi:4506674 /UG=Hs.2280 ribophorin I /FL=gb:NM_002950.1	NM_002950		NP_002941

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201012_at	0.006722	gb:NM_000700.1 /DEF=Homo sapiens annexin A1 (ANXA1), mRNA. /FEA=mRNA /GEN=ANXA1 /PROD=annexin I /DB_XREF=gi:4502100 /UG=Hs.78225 annexin A1 /FL=gb:BC001275.1 gb:NM_000700.1	NM_000700		NP_000691
201017_at	0.013678	eukaryotic translation initiation factor 1A	BE542684	Hs.4310	NP_001403
201019_s_at	0.013803	gb:NM_001412.1 /DEF=Homo sapiens eukaryotic translation initiation factor 1A (EIF1A), mRNA. /FEA=mRNA /GEN=EIF1A /PROD=eukaryotic translation initiation factor 1A /DB_XREF=gi:4503498 /UG=Hs.4310 eukaryotic translation initiation factor 1A /FL=gb:BC000793.1 gb:L18960.1 gb:NM_001412.1	NM_001412		NP_001403
201020_at	7.25E-04	gb:NM_003405.1 /DEF=Homo sapiens tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, eta polypeptide (YWHAH), mRNA. /FEA=mRNA /GEN=YWHAH /PROD=tyrosine 3-monooxygenasetryptophan5-monooxygenase activation protein, eta polypeptide /DB_XREF=gi:4507950 /UG=Hs.75544 tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, eta polypeptide /FL=gb:BC003047.1 gb:L20422.1 gb:NM_003405.1	NM_003405		NP_003396
201038_s_at	9.85E-04	ESTs, Highly similar to putative human HLA class II associated protein I; cerebellar leucine rich acidic nuclear protein [Homo sapiens] [H.sapiens]	BE560202	Hs.356089	

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
201047_x_at	7.76E-04	gb:BC003617.1 /DEF=Homo sapiens, RAB6, member RAS oncogene family, clone MGC:1654, mRNA, complete cds. /FEA=mRNA /PROD=RAB6, member RAS oncogene family /DB_XREF=gi:13177663 /UG=Hs.5636 RAB6A, member RAS oncogene family /FL=gb:BC003617.1 gb:NM_002869.1 gb:AF130986.1 gb:AF130122.1 gb:M28212.1 gb:AF119836.1 gb:AF198616.1	BC003617		NP_002860
201049_s_at	0.047555	gb:NM_022551.1 /DEF=Homo sapiens ribosomal protein S18 (RPS18), mRNA. /FEA=mRNA /GEN=RPS18 /PROD=ribosomal protein S18 /DB_XREF=gi:11968181 /UG=Hs.275865 ribosomal protein S18 /FL=gb:NM_022551.1	NM_022551		NP_072045
201051_at	0.001942	ESTs, Highly similar to putative human HLA class II associated protein I; cerebellar leucine rich acidic nuclear protein [Homo sapiens] [H.sapiens]	BE560202	Hs.356089	
201052_s_at	7.63E-04	proteasome (prosome, macropain) inhibitor subunit 1 (PI31)	BG029917	Hs.75925	NP_848694
201053_s_at	0.00847	gb:NM_006814.1 /DEF=Homo sapiens proteasome (prosome, macropain) inhibitor subunit 1 (PI31) (PSMF1), mRNA. /FEA=mRNA /GEN=PSMF1 /PROD=proteasome inhibitor /DB_XREF=gi:5803122 /UG=Hs.75925 proteasome (prosome, macropain) inhibitor subunit 1 (PI31) /FL=gb:D88378.1 gb:NM_006814.1	NM_006814		NP_848694

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201056_at	0.032758	golgi autoantigen, golgin subfamily b, macrogolgin (with transmembrane signal), 1	N53479	Hs.7844	
201060_x_at	0.001763	guanine nucleotide binding protein 10	A1537887	Hs.160483	NP_004090
201061_s_at	7.63E-04	gb:M81635.1 /DEF=Homo sapiens erythrocyte membrane protein mRNA, complete cds. /FEA=mRNA /GEN=stomatin peptide /PROD=stomatin peptide /DB_XREF=gi:181183 /UG=Hs.160483 erythrocyte membrane protein band 7.2 (stomatin) /FL=gb:M81635.1 gb:NM_004099.1	M81635		NP_004090
201065_s_at	0.020369	gb:NM_001518.1 /DEF=Homo sapiens general transcription factor II, i (GTF2I), mRNA. /FEA=mRNA /GEN=GTF2I /PROD=general transcription factor II, i /DB_XREF=gi:4504202 /UG=Hs.278589 general transcription factor II, i /FL=gb:U77948.1 gb:AF015553.1 gb:AF038969.1 gb:NM_001518.1	NM_001518		NP_127496
201069_at	0.038584	gb:NM_004530.1 /DEF=Homo sapiens matrix metalloproteinase 2 (gelatinase A, 72kD gelatinase, 72kD type IV collagenase) (MMP2), mRNA. /FEA=mRNA /GEN=MMP2 /PROD=matrix metalloproteinase 2 preproprotein /DB_XREF=gi:11342665 /UG=Hs.111301 matrix metalloproteinase 2 (gelatinase A, 72kD gelatinase, 72kD type IV collagenase) /FL=gb:NM_004530.1 gb:BC002576.1	NM_004530		NP_004521
201070_x_at	0.014875	splicing factor 3b, subunit 1, 155kDa	A1739389	Hs.334826	NP_036565

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201076_at	0.048412	gb:NM_005008.1 /DEF=Homo sapiens non-histone chromosome protein 2 (S. cerevisiae)-like 1 (NHP2L1), mRNA. /FEA=mRNA /GEN=NHP2L1 /PROD=non-histone chromosome protein 2 (S.cerevisiae)-like 1 /DB_XREF=gi:4826859 /UG=Hs.182255 non-histone chromosome protein 2 (S. cerevisiae)-like 1 /FL=gb:BC005358.1 gb:D50420.1 gb:AF091076.1 gb:NM_005008.1 gb:AF155235.1	NM_005008		NP_004999
201078_at	6.66E-04	gb:NM_004800.1 /DEF=Homo sapiens transmembrane 9 superfamily member 2 (TM9SF2), mRNA. /FEA=mRNA /GEN=TM9SF2 /PROD=transmembrane 9 superfamily member 2 /DB_XREF=gi:4758873 /UG=Hs.28757 transmembrane 9 superfamily member 2 /FL=gb:U81006.1 gb:NM_004800.1	NM_004800		NP_004791
201086_x_at	0.003965	gb:NM_003103.1 /DEF=Homo sapiens SON DNA binding protein (SON), mRNA. /FEA=mRNA /GEN=SON /PROD=SON DNA binding protein /DB_XREF=gi:4507152 /UG=Hs.92909 SON DNA binding protein /FL=gb:NM_003103.1	NM_003103		NP_620305

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201088_at	0.007839	gb:NM_002266.1 /DEF=Homo sapiens karyopherin alpha 2 (RAG cohort 1, importin alpha 1) (KPNA2), mRNA. /FEA=mRNA /GEN=KPNA2 /PROD=karyopherin alpha 2 /DB_XREF=gi:4504896 /UG=Hs.159557 karyopherin alpha 2 (RAG cohort 1, importin alpha 1) /FL=gb:NM_002266.1 gb:U09559.1 gb:U28386.1	NM_002266		NP_002257
201089_at	0.006596	gb:NM_001693.1 /DEF=Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta polypeptide, 5658kD, isoform 2 (ATP6B2), mRNA. /FEA=mRNA /GEN=ATP6B2 /PROD=ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta polypeptide, 5658kD, isoform 2 /DB_XREF=gi:4502310 /UG=Hs.1697 ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta polypeptide, 5658kD, isoform 2 /FL=gb:BC003100.1 gb:NM_001693.1 gb:L35249.1	NM_001693		NP_001684
201090_x_at	0.001406	gb:NM_006082.1 /DEF=Homo sapiens tubulin, alpha, ubiquitous (K-ALPHA-1), mRNA. /FEA=mRNA /GEN=K-ALPHA-1 /PROD=tubulin, alpha, ubiquitous /DB_XREF=gi:5174476 /UG=Hs.278242 tubulin, alpha, ubiquitous /FL=gb:BC000696.1 gb:BC001128.1 gb:BC001209.1 gb:K00558.1 gb:AF081484.1 gb:NM_006082.1	NM_006082		NP_006073
201091_s_at	0.001763	chromobox homolog 3 (HP1 gamma homolog, Drosophila)	BE748755	Hs.278554	NP_057671

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201093_x_at	0.027219	gb:NM_004168.1 /DEF=Homo sapiens succinate dehydrogenase complex, subunit A, flavoprotein (Fp) (SDHA), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=SDHA /PROD=succinate dehydrogenase complex, subunit A, flavoprotein precursor /DB_XREF=gi:4759079 /UG=Hs.469 succinate dehydrogenase complex, subunit A, flavoprotein (Fp) /FL=gb:BC001380.1 gb:L21936.1 gb:NM_004168.1 gb:D30648.1	NM_004168		NP_004159
201095_at	0.0081	gb:NM_004394.1 /DEF=Homo sapiens death-associated protein (DAP), mRNA. /FEA=mRNA /GEN=DAP /PROD=death-associated protein /DB_XREF=gi:4758119 /UG=Hs.75189 death-associated protein /FL=gb:BC002726.1 gb:NM_004394.1	NM_004394		NP_004385
201096_s_at	0.001179	ADP-ribosylation factor 4	AL537042	Hs.75290	NP_001651
201097_s_at	0.001812	gb:NM_001660.2 /DEF=Homo sapiens ADP-ribosylation factor 4 (ARF4), mRNA. /FEA=mRNA /GEN=ARF4 /PROD=ADP-ribosylation factor 4 /DB_XREF=gi:6995998 /UG=Hs.75290 ADP-ribosylation factor 4 /FL=gb:BC003364.1 gb:M36341.1 gb:NM_001660.2	NM_001660		NP_001651

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201098_at	7.25E-04	gb:NM_004766.1 /DEF=Homo sapiens coatomer protein complex, subunit beta 2 (beta prime) (COPB2), mRNA. /FEA=mRNA /GEN=COPB2 /PROD=coatomer protein complex, subunit beta 2 (betaprime) /DB_XREF=gi:4758031 /UG=Hs.75724 coatomer protein complex, subunit beta 2 (beta prime) /FL=gb:BC000326.1 gb:NM_004766.1	NM_004766		NP_004757
201099_at	6.66E-04	ubiquitin specific protease 9, X chromosome (fat facets-like Drosophila)	AA824386	Hs.77578	
201100_s_at	0.002214	gb:NM_004652.2 /DEF=Homo sapiens ubiquitin specific protease 9, X chromosome (Drosophila fat facets related) (USP9X), transcript variant 1, mRNA. /FEA=mRNA /GEN=USP9X /PROD=Drosophila fat facets related, X-linked, isoform1 /DB_XREF=gi:11641424 /UG=Hs.77578 ubiquitin specific protease 9, X chromosome (Drosophila fat facets related) /FL=gb:NM_004652.2	NM_004652		NP_068706
201102_s_at	0.010171	gb:NM_002626.1 /DEF=Homo sapiens phosphofructokinase, liver (PFKL), mRNA. /FEA=mRNA /GEN=PFKL /PROD=phosphofructokinase, liver /DB_XREF=gi:4505746 /UG=Hs.155455 phosphofructokinase, liver /FL=gb:BC004920.1 gb:X15573.1 gb:NM_002626.1	NM_002626		NP_002617
201103_x_at	0.049833	hypothetical protein DJ328E19.C1.1	BE299495	Hs.323463	NP_775909

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
201104_x_at	0.022707	gb:NM_015383.1 /DEF=Homo sapiens hypothetical protein (DJ328E19.C1.1), mRNA. /FEA=mRNA /GEN=DJ328E19.C1.1 /PROD=hypothetical protein /DB_XREF=gi:7657016 /UG=Hs.218329 hypothetical protein /FL=gb:NM_015383.1	NM_015383		NP_056198
201106_at	0.018398	gb:NM_002085.1 /DEF=Homo sapiens glutathione peroxidase 4 (phospholipid hydroperoxidase) (GPX4), mRNA. /FEA=mRNA /GEN=GPX4 /PROD=glutathione peroxidase 4 /DB_XREF=gi:4504106 /UG=Hs.2706 glutathione peroxidase 4 (phospholipid hydroperoxidase) /FL=gb:NM_002085.1	NM_002085		NP_002076
201114_x_at	0.004987	gb:NM_002792.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA. /FEA=mRNA /GEN=PSMA7 /PROD=proteasome (prosome, macropain) subunit, alpha type, 7 /DB_XREF=gi:4506188 /UG=Hs.233952 proteasome (prosome, macropain) subunit, alpha type, 7 /FL=gb:BC004427.1 gb:AF022815.1 gb:AF054185.1 gb:NM_002792.1	NM_002792		NP_689468

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
201119_s_at	0.013527	gb:NM_004074.1 /DEF=Homo sapiens cytochrome c oxidase subunit VIII (COX8), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=COX8 /PROD=cytochrome c oxidase subunit VIII /DB_XREF=gi:4758043. /UG=Hs.81097 cytochrome c oxidase subunit VIII /FL=gb:NM_004074.1	NM_004074		NP_004065
201128_s_at	0.007818	gb:NM_001096.1 /DEF=Homo sapiens ATP citrate lyase (ACLY), mRNA. /FEA=mRNA /GEN=ACLY /PROD=ATP citrate lyase /DB_XREF=gi:4501864 /UG=Hs.174140 ATP citrate lyase /FL=gb:NM_001096.1	NM_001096		NP_001087
201129_at	0.016463	gb:NM_006276.2 /DEF=Homo sapiens splicing factor, arginineserine-rich 7 (35kD) (SFRS7), mRNA. /FEA=mRNA /GEN=SFRS7 /PROD=splicing factor, arginineserine-rich 7 (35kD) /DB_XREF=gi:6857827 /UG=Hs.184167 splicing factor, arginineserine-rich 7 (35kD) /FL=gb:BC000997.2 gb:L22253.1 gb:NM_006276.2	NM_006276		NP_006267
201132_at	6.66E-04	gb:NM_019597.1 /DEF=Homo sapiens heterogeneous nuclear ribonucleoprotein H2 (H) (HNRPH2), mRNA. /FEA=mRNA /GEN=HNRPH2 /PROD=heterogeneous nuclear ribonucleoprotein H2 (H) /DB_XREF=gi:9624997 /UG=Hs.278857 heterogeneous nuclear ribonucleoprotein H2 (H) /FL=gb:NM_019597.1	NM_019597		NP_062543
201133_s_at	0.004987	KIAA0438 gene product	AA142966	Hs.279849	NP_055634

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	D scription	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
201136_at	0.002749	gb:NM_002668.1 /DEF=Homo sapiens proteolipid protein 2 (colonic epithelium-enriched) (PLP2), mRNA. /FEA=mRNA /GEN=PLP2 /PROD=proteolipid protein 2 (colonic epithelium-enriched) /DB_XREF=gi:4505892 /UG=Hs.77422 proteolipid protein 2 (colonic epithelium-enriched) /FL=gb:L09604.1 gb:NM_002668.1	NM_002668		NP_002659
201138_s_at	0.015926	Sjogren syndrome antigen B (autoantigen La)	BG532929	Hs.83715	NP_003133
201140_s_at	0.031624	gb:NM_004583.1 /DEF=Homo sapiens RAB5C, member RAS oncogene family (RAB5C), mRNA. /FEA=mRNA /GEN=RAB5C /PROD=RAB5C, member RAS oncogene family /DB_XREF=gi:4759019 /UG=Hs.479 RAB5C, member RAS oncogene family /FL=gb:NM_004583.1 gb:U11293.1 gb:U18420.1 gb:AF141304.1	NM_004583		NP_004574
201146_at	7.63E-04	gb:NM_006164.1 /DEF=Homo sapiens nuclear factor (erythroid-derived 2)-like 2 (NFE2L2), mRNA. /FEA=mRNA /GEN=NFE2L2 /PROD=nuclear factor (erythroid-derived 2)-like 2 /DB_XREF=gi:5453775 /UG=Hs.155396 nuclear factor (erythroid-derived 2)-like 2 /FL=gb:NM_006164.1	NM_006164		NP_006155
201154_x_at	0.014936	gb:NM_000968.1 /DEF=Homo sapiens ribosomal protein L4 (RPL4), mRNA. /FEA=mRNA /GEN=RPL4 /PROD=ribosomal protein L4 /DB_XREF=gi:4506652 /UG=Hs.286 ribosomal protein L4 /FL=gb:BC001365.1 gb:L20868.1 gb:D23660.1 gb:NM_000968.1	NM_000968		NP_000959

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201155_s_at	0.049392	gb:NM_014874.1 /DEF=Homo sapiens KIAA0214 gene product (KIAA0214), mRNA. /FEA=mRNA /GEN=KIAA0214 /PROD=KIAA0214 gene product /DB_XREF=gi:7662003 /UG=Hs.3363 KIAA0214 gene product /FL=gb:D86987.1 gb:AF036536.1 gb:NM_014874.1	NM_014874		NP_055689
201160_s_at	7.63E-04	cold shock domain protein A	AL556190	Hs.198726	NP_003642
201161_s_at	8.11E-04	gb:NM_003651.1 /DEF=Homo sapiens cold shock domain protein A (CSDA), mRNA. /FEA=mRNA /GEN=CSDA /PROD=cold shock domain protein A /DB_XREF=gi:4503070 /UG=Hs.1139 cold shock domain protein A /FL=gb:NM_003651.1	NM_003651		NP_003642
201163_s_at	0.001812	gb:NM_001553.1 /DEF=Homo sapiens insulin-like growth factor binding protein 7 (IGFBP7), mRNA. /FEA=mRNA /GEN=IGFBP7 /PROD=insulin-like growth factor binding protein 7 /DB_XREF=gi:4504618 /UG=Hs.119206 insulin-like growth factor binding protein 7 /FL=gb:L19182.1 gb:NM_001553.1	NM_001553		NP_001544
201165_s_at	7.25E-04	pumilio homolog 1 (Drosophila)	BG474429	Hs.153834	
201166_s_at	0.01135	gb:NM_014676.1 /DEF=Homo sapiens pumilio (Drosophila) homolog 1 (PUM1), mRNA. /FEA=mRNA /GEN=PUM1 /PROD=pumilio (Drosophila) homolog 1 /DB_XREF=gi:13491165 /UG=Hs.153834 pumilio (Drosophila) homolog 1 /FL=gb:AF315592.1 gb:NM_014676.1	NM_014676		NP_055491

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201170_s_at	0.011166	gb:NM_003670.1 /DEF=Homo sapiens basic helix-loop-helix domain containing, class B, 2 (BHLHB2), mRNA. /FEA=mRNA /GEN=BHLHB2 /PROD=differentiated embryo chondrocyte expressed gene1 /DB_XREF=gi:4503298 /UG=Hs.171825 basic helix-loop-helix domain containing, class B, 2 /FL=gb:AB004066.1 gb:NM_003670.1	NM_003670		NP_003661
201171_at	0.030518	Consensus includes gb:NM_003945.1 /DEF=Homo sapiens ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 9kD (ATP6H), mRNA. /FEA=mRNA /GEN=ATP6H /PROD=ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 9kD /DB_XREF=gi:4502318 /UG=Hs.24322 ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 9kD /FL=gb:NM_003945.1	NM_003945		NP_003936
201172_x_at	0.003875	gb:NM_003945.1 /DEF=Homo sapiens ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 9kD (ATP6H), mRNA. /FEA=mRNA /GEN=ATP6H /PROD=ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 9kD /DB_XREF=gi:4502318 /UG=Hs.24322 ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 9kD /FL=gb:NM_003945.1	NM_003945		NP_003936

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
201173_x_at	0.003373	gb:NM_006600.1 /DEF=Homo sapiens nuclear distribution gene C (A.nidulans) homolog (NUDC), mRNA. /FEA=mRNA /GEN=NUDC /PROD=nuclear distribution gene C (A.nidulans)homolog /DB_XREF=gi:5729952 /UG=Hs.263812 nuclear distribution gene C (A.nidulans) homolog /FL=gb:BC002399.1 gb:BC003132.1 gb:AB019408.1 gb:AF130736.1 gb:AF125465.1 gb:AF100760.1 gb:NM_006600.1	NM_006600		NP_006591
201174_s_at	0.002446	gb:NM_018975.1 /DEF=Homo sapiens TRF2-interacting telomeric RAP1 protein (RAP1), mRNA. /FEA=mRNA /GEN=RAP1 /PROD=TRF2-interacting telomeric RAP1 protein /DB_XREF=gi:9507032 /UG=Hs.274428 TRF2-interacting telomeric RAP1 protein /FL=gb:BC004465.1 gb:AF262988.1 gb:NM_018975.1	NM_018975		NP_061848
201176_s_at	0.006596	gb:NM_001655.2 /DEF=Homo sapiens archain 1 (ARCN1), mRNA. /FEA=mRNA /GEN=ARCN1 /PROD=archain /DB_XREF=gi:11863153 /UG=Hs.33642 archain 1 /FL=gb:NM_001655.2	NM_001655		NP_001646
201178_at	7.63E-04	gb:NM_012179.1 /DEF=Homo sapiens F-box only protein 7 (FBXO7), mRNA. /FEA=mRNA /GEN=FBXO7 /PROD=F-box only protein 7 /DB_XREF=gi:7106310 /UG=Hs.5912 F-box only protein 7 /FL=gb:AF129537.1 gb:NM_012179.1 gb:AF233225.1	NM_012179		NP_036311

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201179_s_at	0.010277	gb:J03005.1 /DEF=Human alternative guanine nucleotide-binding regulatory protein (G) alpha-inhibitory-subunit mRNA, complete cds. /FEA=mRNA /GEN=GNAI1 /DB_XREF=gi:183183 /UG=Hs.73799 guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3 /FL=gb:J03005.1 gb:J03198.1 gb:M27543.1 gb:J03238.1 gb:NM_006496.1	J03005		NP_006487
201180_s_at	6.66E-04	gb:J03198.1 /DEF=Human stimulatory G protein (of receptor-regulated K ⁺ channels) alpha subunit mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:183224 /UG=Hs.73799 guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3 /FL=gb:J03005.1 gb:J03198.1 gb:M27543.1 gb:J03238.1 gb:NM_006496.1	J03198		NP_006487
201181_at	0.005652	gb:NM_006496.1 /DEF=Homo sapiens guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3 (GNAI3), mRNA. /FEA=mRNA /GEN=GNAI3 /PROD=guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3 /DB_XREF=gi:5729849 /UG=Hs.73799 guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3 /FL=gb:J03005.1 gb:J03198.1 gb:M27543.1 gb:J03238.1 gb:NM_006496.1	NM_006496		NP_006487
201183_s_at	0.016179	chromodomain helicase DNA binding protein 4	AI761771	Hs.74441	NP_001264

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201186_at	0.001719	gb:NM_002337.1 /DEF=Homo sapiens low density lipoprotein-related protein-associated protein 1 (alpha-2-macroglobulin receptor-associated protein 1) (LRPAP1), mRNA. /FEA=mRNA /GEN=LRPAP1 /PROD=low density lipoprotein-related protein-associated protein 1 (alpha-2-macroglobulin receptor-associated protein 1) /DB_XREF=gi:4505020 /UG=Hs.75140 low density lipoprotein-related protein-associated protein 1 (alpha-2-macroglobulin receptor-associated protein 1) /FL=gb:M63959.1 gb:NM_002337.1	NM_002337		NP_002328
201190_s_at	0.010713	phosphatidylinositol transfer protein	H15647	Hs.409367	NP_006215
201191_at	0.004483	phosphatidylinositol transfer protein	H15647	Hs.409367	NP_006215
201192_s_at	9.85E-04	gb:NM_006224.1 /DEF=Homo sapiens phosphatidylinositol transfer protein (PITPN), mRNA. /FEA=mRNA /GEN=PITPN /PROD=phosphatidylinositol transfer protein /DB_XREF=gi:5453907 /UG=Hs.79709 phosphatidylinositol transfer protein /FL=gb:D30036.1 gb:M73704.1 gb:NM_006224.1	NM_006224		NP_006215

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gen Acc sion No.	Unigene Accession No.	Protein Accession No.
201193_at	7.63E-04	gb:NM_005896.1 /DEF=Homo sapiens isocitrate dehydrogenase 1 (NADP+), soluble (IDH1), mRNA. /FEA=mRNA /GEN=IDH1 /PROD=isocitrate dehydrogenase 1 (NADP+), soluble /DB_XREF=gi:5174470 /UG=Hs.11223 isocitrate dehydrogenase 1 (NADP+), soluble /FL=gb:AF020038.1 gb:AF113917.1 gb:NM_005896.1 gb:AL136702.1	NM_005896		NP_005887
201194_at	0.039362	gb:NM_003009.1 /DEF=Homo sapiens selenoprotein W, 1 (SEPW1), mRNA. /FEA=mRNA /GEN=SEPW1 /PROD=selenoprotein W, 1 /DB_XREF=gi:4506886 /UG=Hs.14231 selenoprotein W, 1 /FL=gb:U67171.1 gb:AF015283.1 gb:NM_003009.1	NM_003009		NP_003000
201196_s_at	0.002109	gb:M21154.1 /DEF=Human S-adenosylmethionine decarboxylase mRNA, complete cds. /FEA=mRNA /GEN=AMD2 /DB_XREF=gi:178517 /UG=Hs.262476 S-adenosylmethionine decarboxylase 1 /FL=gb:BC000171.2 gb:M21154.1 gb:NM_001634.3	M21154		NP_001625
201200_at	0.001763	gb:NM_003851.1 /DEF=Homo sapiens cellular repressor of E1A-stimulated genes (CREG), mRNA. /FEA=mRNA /GEN=CREG /PROD=cellular repressor of E1A-stimulated genes /DB_XREF=gi:4503036 /UG=Hs.5710 cellular repressor of E1A-stimulated genes /FL=gb:AF084523.1 gb:NM_003851.1	NM_003851		NP_003842

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201201_at	0.038202	gb:NM_000100.1 /DEF=Homo sapiens cystatin B (stefin B) (CSTB), mRNA. /FEA=mRNA /GEN=CSTB /PROD=cystatin B (stefin B) /DB_XREF=gi:4503116 /UG=Hs.695 cystatin B (stefin B) /FL=gb:BC003370.1 gb:L03558.1 gb:NM_000100.1	NM_000100		NP_000091
201202_at	0.002214	gb:NM_002592.1 /DEF=Homo sapiens proliferating cell nuclear antigen (PCNA), mRNA. /FEA=mRNA /GEN=PCNA /PROD=proliferating cell nuclear antigen /DB_XREF=gi:4505640 /UG=Hs.78996 proliferating cell nuclear antigen /FL=gb:BC000491.1 gb:M15796.1 gb:NM_002592.1	NM_002592		NP_002583
201214_s_at	0.02217	gb:NM_002712.1 /DEF=Homo sapiens protein phosphatase 1, regulatory subunit 7 (PPP1R7), mRNA. /FEA=mRNA /GEN=PPP1R7 /PROD=protein phosphatase 1, regulatory subunit 7 /DB_XREF=gi:4506012 /UG=Hs.36587 protein phosphatase 1, regulatory subunit 7 /FL=gb:BC000910.1 gb:NM_002712.1	NM_002712		NP_002703
201218_at	0.01747	Consensus includes gb:N23018 /FEA=EST /DB_XREF=gi:1137168 /DB_XREF=est:yx65d12.s1 /CLONE=IMAGE:266615 /UG=Hs.171391 C-terminal binding protein 2 /FL=gb:AF016507.1 gb:NM_001329.1	NM_001329		NP_073713

Genes Corresponding To Differentially Expressed Genes in Figur 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
201221_s_at	0.003842	gb:NM_003089.1 /DEF=Homo sapiens small nuclear ribonucleoprotein 70kD polypeptide (RNP antigen) (SNRP70), mRNA. /FEA=mRNA /GEN=SNRP70 /PROD=small nuclear ribonucleoprotein 70kD polypeptide(RNP antigen) /DB_XREF=gi:4507118 /UG=Hs.174051 small nuclear ribonucleoprotein 70kD polypeptide (RNP antigen) /FL=gb:BC000342.1 gb:M22636.1 gb:NM_003089.1	NM_003089		NP_003080
201222_s_at	7.63E-04	RAD23 homolog B (S. cerevisiae)	AL527365	Hs.178658	NP_002865
201223_s_at	0.002621	gb:NM_002874.1 /DEF=Homo sapiens RAD23 (S. cerevisiae) homolog B (RAD23B), mRNA. /FEA=mRNA /GEN=RAD23B /PROD=RAD23 (S. cerevisiae) homolog B /DB_XREF=gi:4506386 /UG=Hs.178658 RAD23 (S. cerevisiae) homolog B /FL=gb:NM_002874.1 gb:D21090.1	NM_002874		NP_002865
201234_at	0.003315	gb:NM_004517.1 /DEF=Homo sapiens integrin-linked kinase (ILK), mRNA. /FEA=mRNA /GEN=ILK /PROD=integrin-linked kinase /DB_XREF=gi:4758605 /UG=Hs.6196 integrin-linked kinase /FL=gb:U40282.1 gb:NM_004517.1	NM_004517		NP_004508
201237_at	0.003669	capping protein (actin filament) muscle Z-line, alpha 2	AV685920	Hs.75546	NP_006127

Genes Corrsponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
201238_s_at	0.001167	gb:BC005338.1 /DEF=Homo sapiens, capping protein (actin filament) muscle Z-line, alpha 2, clone MGC:12426, mRNA, complete cds. /FEA=mRNA /PROD=capping protein (actin filament) muscle Z-line, alpha 2 /DB_XREF=gi:13529130 /UG=Hs.75546 capping protein (actin filament) muscle Z-line, alpha 2 /FL=gb:BC005338.1 gb:NM_006136.1 gb:U03269.1	BC005338		NP_006127
201239_s_at	0.011487	ESTs, Highly similar to SP25_HUMAN Microsomal signal peptidase 25 kDa subunit (SPase 25 kDa subunit) (SPC25) [H.sapiens]	BF530535	Hs.381155	
201248_s_at	0.037297	gb:NM_004599.1 /DEF=Homo sapiens sterol regulatory element binding transcription factor 2 (SREBF2), mRNA. /FEA=mRNA /GEN=SREBF2 /PROD=sterol regulatory element binding transcriptionfactor 2 /DB_XREF=gi:4759169 /UG=Hs.108689 sterol regulatory element binding transcription factor 2 /FL=gb:U02031.1 gb:NM_004599.1	NM_004599		NP_004590
201255_x_at	0.038202	gb:NM_004639.1 /DEF=Homo sapiens HLA-B associated transcript-3 (D6S52E), mRNA. /FEA=mRNA /GEN=D6S52E /PROD=HLA-B associated transcript-3 /DB_XREF=gi:4758109 /UG=Hs.274348 HLA-B associated transcript-3 /FL=gb:M33519.1 gb:NM_004639.1	NM_004639		NP_542434

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unig. n Accession No.	Protein Accession No.
201256_at	0.001179	gb:NM_004718.1 /DEF=Homo sapiens cytochrome c oxidase subunit VIIa polypeptide 2 like (COX7A2L), mRNA. /FEA=mRNA /GEN=COX7A2L /PROD=cytochrome c oxidase subunit VIIa polypeptide 2like /DB_XREF=gi:4758041 /UG=Hs.30888 cytochrome c oxidase subunit VIIa polypeptide 2 like /FL=gb:BC005251.1 gb:AB007618.1 gb:NM_004718.1	NM_004718		NP_004709
201264_at	0.003674	gb:NM_007263.1 /DEF=Homo sapiens coatomer protein complex, subunit epsilon (COPE), mRNA. /FEA=mRNA /GEN=COPE /PROD=coatomer protein complex, subunit epsilon /DB_XREF=gi:6005734 /UG=Hs.10326 coatomer protein complex, subunit epsilon /FL=gb:AL136928.1 gb:BC003155.1 gb:NM_007263.1	NM_007263		NP_009194
201266_at	7.63E-04	gb:NM_003330.1 /DEF=Homo sapiens thioredoxin reductase 1 (TXNRD1), mRNA. /FEA=mRNA /GEN=TXNRD1 /PROD=thioredoxin reductase 1 /DB_XREF=gi:4507746 /UG=Hs.13046 thioredoxin reductase 1 /FL=gb:D88687.1 gb:AF077367.1 gb:NM_003330.1 gb:AF208018.1	NM_003330		NP_003321

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201273_s_at	0.002559	gb:NM_003133.1 /DEF=Homo sapiens signal recognition particle 9kD (SRP9), mRNA. /FEA=mRNA /GEN=SRP9 /PROD=signal recognition particle 9kD. /DB_XREF=gi:4507216. /UG=Hs.75975 signal recognition particle 9kD /FL=gb:NM_003133.1 gb:U20998.1	NM_003133		NP_003124
201275_at	0.001108	gb:NM_002004.1 /DEF=Homo sapiens farnesyl diphosphate synthase (farnesyl pyrophosphate synthetase, dimethylallyltranstransferase, geranyltranstransferase) (FDPS), mRNA. /FEA=mRNA /GEN=FDPS /PROD=farnesyl diphosphate synthase (farnesylpyrophosphate synthetase, dimethylallyltranstransferase, geranyltranstransferase) /DB_XREF=gi:4503684 /UG=Hs.77393 farnesyl diphosphate synthase (farnesyl pyrophosphate synthetase, dimethylallyltranstransferase, geranyltranstransferase) /FL=gb:J05262.1 gb:D14697.1 gb:NM_002004.1	NM_002004		NP_001995
201277_s_at	0.001719	gb:NM_004499.1 /DEF=Homo sapiens heterogeneous nuclear ribonucleoprotein AB (HNRPAB), mRNA. /FEA=mRNA /GEN=HNRPAB /PROD=heterogeneous nuclear ribonucleoprotein AB /DB_XREF=gi:4758541 /UG=Hs.81361 heterogeneous nuclear ribonucleoprotein AB /FL=gb:BC002625.1 gb:BC004561.1 gb:M65028.1 gb:NM_004499.1	NM_004499		NP_112556

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201285_at	7.25E-04	gb:NM_013446.1 /DEF=Homo sapiens makorin, ring finger protein, 1 (MKRN1); mRNA. /FEA=mRNA /GEN=MKRN1 /PROD=makorin, ring finger protein, 1 /DB_XREF=gi:7305272 /UG=Hs.7838 makorin, ring finger protein, 1 /FL=gb:AL136812.1 gb:AF192784.1 gb:NM_013446.1	NM_013446		NP_038474
201290_at	0.016179	gb:NM_014300.1 /DEF=Homo sapiens signal peptidase complex (18kD) (SPC18); mRNA. /FEA=mRNA /GEN=SPC18 /PROD=signal peptidase complex (18kD) /DB_XREF=gi:7657608 /UG=Hs.9534 signal peptidase complex (18kD) /FL=gb:BC000359.1 gb:AF061737.1 gb:AF108945.1 gb:NM_014300.1	NM_014300		NP_055115
201296_s_at	0.010065	gb:NM_015626.1 /DEF=Homo sapiens DKFZP564A122 protein (DKFZP564A122); mRNA. /FEA=mRNA. /GEN=DKFZP564A122 /PROD=DKFZP564A122 protein /DB_XREF=gi:7661595 /UG=Hs.187991 DKFZP564A122 protein /FL=gb:AF106684.1 gb:NM_015626.1	NM_015626		NP_599027
201297_s_at	0.009378	Consensus includes gb:AK023321.1 /DEF=Homo sapiens cDNA FLJ13259 fis, clone OVARC1000876, moderately similar to MOB1 PROTEIN. /FEA=mRNA /DB_XREF=gi:10435206 /UG=Hs.196437 hypothetical protein FLJ10788 /FL=gb:AB016839.1 gb:BC003398.1 gb:NM_018221.1	NM_018221		NP_060691

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201298_s_at	0.00109	gb:BC003398.1 /DEF=Homo sapiens, hypothetical protein FLJ10788, clone MGC:4929, mRNA, complete cds. /FEA=mRNA /PROD=hypothetical protein FLJ10788 /DB_XREF=gi:13097287 /UG=Hs.196437 hypothetical protein FLJ10788 /FL=gb:AB016839.1 gb:BC003398.1 gb:NM_018221.1	BC003398		NP_060691
201301_s_at	7.25E-04	gb:BC000182.1 /DEF=Homo sapiens, annexin A4, clone MGC:2271, mRNA, complete cds. /FEA=mRNA /PROD=annexin A4 /DB_XREF=gi:12652858 /UG=Hs.77840 annexin A4 /FL=gb:D78152.1 gb:BC000182.1 gb:M82809.1 gb:M19383.1 gb:NM_001153.2	BC000182		NP_001144
201302_at	6.66E-04	gb:NM_001153.2 /DEF=Homo sapiens annexin A4 (ANXA4), mRNA. /FEA=mRNA /GEN=ANXA4 /PROD=annexin IV /DB_XREF=gi:4809272 /UG=Hs.77840 annexin A4 /FL=gb:D78152.1 gb:BC000182.1 gb:M82809.1 gb:M19383.1 gb:NM_001153.2	NM_001153		NP_001144
201303_at	0.001167	gb:NM_014740.1 /DEF=Homo sapiens KIAA0111 gene product (KIAA0111), mRNA. /FEA=mRNA /GEN=KIAA0111 /PROD=KIAA0111 gene product /DB_XREF=gi:7661919 /UG=Hs.79768 KIAA0111 gene product /FL=gb:BC003662.1 gb:BC004386.1 gb:D21853.1 gb:NM_014740.1	NM_014740		NP_055555

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCanc r					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201311_s_at	6.66E-04	SH3 domain binding glutamic acid-rich protein like	AL515318	Hs.408289	NP_003013
201312_s_at	9.85E-04	gb:NM_003022.1 /DEF=Homo sapiens SH3 domain binding glutamic acid-rich protein like (SH3BGRL), mRNA. /FEA=mRNA /GEN=SH3BGRL /PROD=SH3 domain binding glutamic acid-rich proteinlike /DB_XREF=gi:4506924 /UG=Hs.14368 SH3 domain binding glutamic acid-rich protein like /FL=gb:AL136718.1 gb:AF042081.1 gb:NM_003022.1	NM_003022		NP_003013
201317_s_at	0.002559	gb:NM_002787.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 2 (PSMA2), mRNA. /FEA=mRNA /GEN=PSMA2 /PROD=proteasome (prosome, macropain) subunit, alphas type, 2 /DB_XREF=gi:4506180 /UG=Hs.181309 proteasome (prosome, macropain) subunit, alpha type, 2 /FL=gb:NM_002787.1	NM_002787		NP_002778
201318_s_at	7.63E-04	gb:NM_006471.1 /DEF=Homo sapiens myosin, light polypeptide, regulatory, non-sarcomeric (20kD) (MLCB), mRNA. /FEA=mRNA /GEN=MLCB /PROD=myosin, light polypeptide, regulatory,non-sarcomeric (20kD) /DB_XREF=gi:5453739 /UG=Hs.233936 myosin, light polypeptide, regulatory, non-sarcomeric (20kD) /FL=gb:NM_006471.1	NM_006471		NP_006462

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201319_at	0.03947	gb:NM_006471.1 /DEF=Homo sapiens myosin, light polypeptide, regulatory, non-sarcomeric (20kD) (MLCB), mRNA. /FEA=mRNA /GEN=MLCB /PROD=myosin, light polypeptide, regulatory, non-sarcomeric (20kD) /DB_XREF=gi:5453739 /UG=Hs.233936 myosin, light polypeptide, regulatory, non-sarcomeric (20kD) /FL=gb:NM_006471.1	NM_006471		NP_006462
201322_at	0.003669	gb:NM_001686.1 /DEF=Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F1 complex, beta polypeptide (ATP5B), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=ATP5B /PROD=ATP synthase, H ⁺ transporting, mitochondrial F1 complex, beta polypeptide /DB_XREF=gi:4502294 /UG=Hs.25 ATP synthase, H ⁺ transporting, mitochondrial F1 complex, beta polypeptide /FL=gb:D00022.1 gb:NM_001686.1	NM_001686		NP_001677
201328_at	9.85E-04	v-ets erythroblastosis virus E26 oncogene homolog 2 (avian)	AL575509	Hs.85146	
201336_at	0.001213	gb:BC003570.1 /DEF=Homo sapiens, Similar to vesicle-associated membrane protein 3, clone MGC:2110, mRNA, complete cds. /FEA=mRNA /PROD=Similar to vesicle-associated membrane protein3 /DB_XREF=gi:13097737 /UG=Hs.66708 vesicle-associated membrane protein 3 (cellubrevin) /FL=gb:BC003570.1 gb:NM_004781.2	BC003570		NP_004772

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201339_s_at	0.030776	gb:NM_002979.1 /DEF=Homo sapiens sterol carrier protein 2 (SCP2), mRNA. /FEA=mRNA /GEN=SCP2 /PROD=sterol carrier protein 2 /DB_XREF=gi:4506822 /UG=Hs.75760 sterol carrier protein 2 /FL=gb:M75883.1 gb:M75884.1 gb:M55421.1 gb:NM_002979.1	NM_002979		NP_002970
201343_at	0.003669	Consensus includes gb:BE621259 /FEA=EST /DB_XREF=gi:9892197 /DB_XREF=est:601493415F1 /CLONE=IMAGE:3895850 /UG=Hs.108332 ubiquitin-conjugating enzyme E2D 2 (homologous to yeast UBC45) /FL=gb:U39317.1 gb:NM_003339.1	NM_003339		NP_003330
201345_s_at	6.66E-04	gb:NM_003339.1 /DEF=Homo sapiens ubiquitin-conjugating enzyme E2D 2 (homologous to yeast UBC45) (UBE2D2), mRNA. /FEA=mRNA /GEN=UBE2D2 /PROD=ubiquitin-conjugating enzyme E2D 2 (homologous to yeast UBC45) /DB_XREF=gi:4507774 /UG=Hs.108332 ubiquitin-conjugating enzyme E2D 2 (homologous to yeast UBC45) /FL=gb:U39317.1 gb:NM_003339.1	NM_003339		NP_003330
201346_at	0.030968	gb:NM_024551.1 /DEF=Homo sapiens hypothetical protein FLJ21432 (FLJ21432), mRNA. /FEA=mRNA /GEN=FLJ21432 /PROD=hypothetical protein FLJ21432 /DB_XREF=gi:13375714 /UG=Hs.11641 hypothetical protein FLJ21432 /FL=gb:NM_024551.1 gb:BC004906.1	NM_024551		NP_078827

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unig n Acc ssion No.	Protein Accession No.
201347_x_at	0.020971	gb:NM_012203.1 /DEF=Homo sapiens glyoxylate reductasehydroxypyruvate reductase (GRHPR), mRNA. /FEA=mRNA /GEN=GRHPR /PROD=glyoxylate reductasehydroxypyruvate reductase /DB_XREF=gi:6912395 /UG=Hs.155742 glyoxylate reductasehydroxypyruvate reductase /FL=gb:AF113215.1 gb:BC000605.1 gb:BC003131.1 gb:AF146018.1 gb:AF113251.1 gb:AF134895.1 gb:NM_012203.1	NM_012203		NP_036335
201351_s_at	0.005652	gb:AF070656.1 /DEF=Homo sapiens FtsH homolog mRNA, complete cds. /FEA=mRNA /PROD=FtsH homolog /DB_XREF=gi:4454687 /UG=Hs.206521 YME1 (S.cerevisiae)-like 1 /FL=gb:AF070656.1 gb:NM_014263.1	AF070656		NP_647474
201352_at	0.001179	gb:NM_014263.1 /DEF=Homo sapiens YME1 (S.cerevisiae)-like 1 (YME1L1), mRNA. /FEA=mRNA /GEN=YME1L1 /PROD=YME1 (S.cerevisiae)-like 1 /DB_XREF=gi:7657688 /UG=Hs.206521 YME1 (S.cerevisiae)-like 1 /FL=gb:AF070656.1 gb:NM_014263.1	NM_014263		NP_647474

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201359_at	0.001852	gb:NM_016451.1 /DEF=Homo sapiens coatamer protein complex, subunit beta (COPB), mRNA. /FEA=mRNA /GEN=COPB /PROD=coatamer protein complex, subunit beta /DB_XREF=gi:7705368 /UG=Hs.3059 coatamer protein complex, subunit beta /FL=gb:AF084457.1 gb:AL136593.1 gb:NM_016451.1	NM_016451		NP_057535
201362_at	0.003976	gb:AF205218.1 /DEF=Homo sapiens NS1-binding protein-like protein mRNA, complete cds. /FEA=mRNA /PROD=NS1 binding protein-like protein /DB_XREF=gi:12003206 /UG=Hs.197298 NS1-binding protein /FL=gb:AF205218.1 gb:AB020657.1 gb:AF161553.1 gb:NM_016389.1	AF205218		NP_006460
201363_s_at	7.25E-04	gb:AB020657.1 /DEF=Homo sapiens mRNA for KIAA0850 protein, complete cds. /FEA=mRNA /GEN=KIAA0850 /PROD=KIAA0850 protein /DB_XREF=gi:4240188 /UG=Hs.197298 NS1-binding protein /FL=gb:AF205218.1 gb:AB020657.1 gb:AF161553.1 gb:NM_016389.1	AB020657		NP_006460

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201369_s_at	0.003584	gb:NM_006887.1 /DEF=Homo sapiens butyrate response factor 2 (EGF-response factor 2) (BRF2), mRNA. /FEA=mRNA /GEN=BRF2 /PROD=butyrate response factor 2 (EGF-response factor2) /DB_XREF=gi:5901899 /UG=Hs.78909 butyrate response factor 2 (EGF-response factor 2) /FL=gb:BC005010.1 gb:NM_006887.1	NM_006887		NP_008818
201371_s_at	0.004519	gb:AF062537.1 /DEF=Homo sapiens cullin 3 mRNA, complete cds. /FEA=mRNA /PROD=cullin 3 /DB_XREF=gi:3139078 /UG=Hs.78946 cullin 3 /FL=gb:AF062537.1 gb:AB014517.1 gb:AF052147.1 gb:AF064087.1 gb:NM_003590.1	AF062537		NP_003581
201375_s_at	0.001179	gb:NM_004156.1 /DEF=Homo sapiens protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform (PPP2CB), mRNA. /FEA=mRNA /GEN=PPP2CB /PROD=protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform /DB_XREF=gi:4758951 /UG=Hs.80350 protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform /FL=gb:NM_004156.1	NM_004156		NP_004147

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201381_x_at	0.029624	gb:AF057356.1 /DEF=Homo sapiens calcyclin binding protein mRNA, complete cds. /FEA=mRNA /PROD=calcyclin binding protein /DB_XREF=gi:3063652 /UG=Hs.27258 calcyclin binding protein /FL=gb:AF314752.1 gb:AF057356.1 gb:NM_014412.1	AF057356		NP_055227
201383_s_at	0.044301	membrane component, chromosome 17, surface marker 2 (ovarian carcinoma antigen CA125)	AL044170	Hs.277721	NP_114068
201384_s_at	0.007693	gb:NM_005899.1 /DEF=Homo sapiens membrane component, chromosome 17, surface marker 2 (ovarian carcinoma antigen CA125) (M17S2), mRNA. /FEA=mRNA /GEN=M17S2 /PROD=membrane component, chromosome 17, surfacemarker 2 (ovarian carcinoma antigen CA125) /DB_XREF=gi:5174504 /UG=Hs.277721 membrane component, chromosome 17, surface marker 2 (ovarian carcinoma antigen CA125) /FL=gb:D30756.1 gb:NM_005899.1	NM_005899		NP_114068
201386_s_at	0.041179	gb:AF279891.1 /DEF=Homo sapiens dead box protein 15 mRNA, complete cds. /FEA=mRNA /PROD=dead box protein 15 /DB_XREF=gi:9624452 /UG=Hs.5683 DEADH (Asp-Glu-Ala-AspHis) box polypeptide 15 /FL=gb:AB001636.1 gb:NM_001358.1 gb:AF279891.1	AF279891		NP_001349

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
201390_s_at	0.005119	gb:NM_001320.1 /DEF=Homo sapiens casein kinase 2, beta polypeptide (CSNK2B), mRNA. /FEA=mRNA /GEN=CSNK2B /PROD=casein kinase 2, beta polypeptide /DB_XREF=gi:10334850 /UG=Hs.165843 casein kinase 2, beta polypeptide /FL=gb:NM_001320.1	NM_001320		NP_001311
201400_at	0.001719	gb:NM_002795.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, beta type, 3 (PSMB3), mRNA. /FEA=mRNA /GEN=PSMB3 /PROD=proteasome (prosome, macropain) subunit, betatype, 3 /DB_XREF=gi:4506196 /UG=Hs.82793 proteasome (prosome, macropain) subunit, beta type, 3 /FL=gb:NM_002795.1 gb:D26598.1	NM_002795		NP_002786
201403_s_at	0.04696	gb:NM_004528.1 /DEF=Homo sapiens microsomal glutathione S-transferase 3 (MGST3), mRNA. /FEA=mRNA /GEN=MGST3 /PROD=microsomal glutathione S-transferase 3 /DB_XREF=gi:4758713 /UG=Hs.111811 microsomal glutathione S-transferase 3 /FL=gb:BC000505.1 gb:BC003034.1 gb:AF026977.1 gb:NM_004528.1	NM_004528		NP_004519
201407_s_at	0.025675	protein phosphatase 1, catalytic subunit, beta isoform	AI186712	Hs.21537	NP_002700
201408_at	0.049282	protein phosphatase 1, catalytic subunit, beta isoform	AI186712	Hs.21537	NP_002700

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201409_s_at	7.25E-04	gb:NM_002709.1 /DEF=Homo sapiens protein phosphatase 1, catalytic subunit, beta isoform (PPP1CB), mRNA. /FEA=mRNA /GEN=PPP1CB /PROD=protein phosphatase 1, catalytic subunit, beta isoform /DB_XREF=gi:4506004 /UG=Hs.21537 protein phosphatase 1, catalytic subunit, beta isoform /FL=gb:NM_002709.1 gb:AF092905.1	NM_002709		NP_002700
201410_at	0.00121	ESTs	AI983043	Hs.409207	
201412_at	0.003009	gb:NM_014045.1 /DEF=Homo sapiens DKFZP564C1940 protein (DKFZP564C1940), mRNA. /FEA=mRNA /GEN=DKFZP564C1940 /PROD=DKFZP564C1940 protein /DB_XREF=gi:13027587 /UG=Hs.3804 DKFZP564C1940 protein /FL=gb:BC000424.1 gb:NM_014045.1 gb:AF131760.1	NM_014045		NP_054764
201422_at	0.003393	gb:NM_006332.1 /DEF=Homo sapiens interferon, gamma-inducible protein 30 (IFI30), mRNA. /FEA=mRNA /GEN=IFI30 /PROD=interferon, gamma-inducible protein 30 /DB_XREF=gi:5453695 /UG=Hs.14623 interferon, gamma-inducible protein 30 /FL=gb:J03909.1 gb:NM_006332.1 gb:AF097362.1	NM_006332		NP_006323
201423_s_at	0.019448	cullin 4A	AL037208	Hs.183874	NP_003580

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201425_at	0.002109	gb:NM_000690.1 /DEF=Homo sapiens aldehyde dehydrogenase 2, mitochondrial (ALDH2), mRNA. /FEA=mRNA /GEN=ALDH2 /PROD=aldehyde dehydrogenase 2, mitochondrial /DB_XREF=gi:4502032 /UG=Hs.195432 aldehyde dehydrogenase 2 family (mitochondrial) /FL=gb:BC002967.1 gb:NM_000690.1	NM_000690		NP_000681
201426_s_at	0.018839	vimentin	AI922599	Hs.297753	NP_000995
201429_s_at	0.020369	gb:NM_000998.1 /DEF=Homo sapiens ribosomal protein L37a (RPL37A), mRNA. /FEA=mRNA /GEN=RPL37A /PROD=ribosomal protein L37a /DB_XREF=gi:4506642 /UG=Hs.5566 ribosomal protein L37a /FL=gb:BC000555.1 gb:L06499.1 gb:NM_000998.1	NM_000998		NP_000989
201433_s_at	0.030776	gb:NM_014754.1 /DEF=Homo sapiens phosphatidylserine synthase 1 (PTDSS1), mRNA. /FEA=mRNA /GEN=PTDSS1 /PROD=phosphatidylserine synthase 1 /DB_XREF=gi:7662646 /UG=Hs.77329 phosphatidylserine synthase 1 /FL=gb:BC004192.1 gb:BC004390.1 gb:D14694.1 gb:NM_014754.1	NM_014754		NP_055569
201435_s_at	0.013199	Consensus includes gb:AW268640 /FEA=EST /DB_XREF=gi:6655670 /DB_XREF=est:xv52a03.x1 /CLONE=IMAGE:2816716 /UG=Hs.79306 eukaryotic translation initiation factor 4E /FL=gb:M15353.1 gb:NM_001968.1	NM_001968		NP_001959

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
201441_at	0.003965	gb:NM_001863.2 /DEF=Homo sapiens cytochrome c oxidase subunit VIb (COX6B), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=COX6B /PROD=cytochrome c oxidase subunit VIb /DB_XREF=gi:6680989 /UG=Hs.174031 cytochrome c oxidase subunit VIb /FL=gb:BC001015.1 gb:BC002478.1 gb:NM_001863.2	NM_001863		NP_001854
201444_s_at	7.25E-04	gb:NM_005765.1 /DEF=Homo sapiens ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) membrane sector associated protein M8-9 (APT6M8-9), mRNA. /FEA=mRNA /GEN=APT6M8-9 /PROD=ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) membrane sector associated protein M8-9 /DB_XREF=gi:5031590 /UG=Hs.183434 ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) membrane sector associated protein M8-9 /FL=gb:AF248966.1 gb:NM_005765.1	NM_005765		NP_005756
201453_x_at	0.002692	gb:NM_005614.1 /DEF=Homo sapiens Ras homolog enriched in brain 2 (RHEB2), mRNA. /FEA=mRNA /GEN=RHEB2 /PROD=Ras homolog enriched in brain 2 /DB_XREF=gi:5032040 /UG=Hs.279903 Ras homolog enriched in brain 2 /FL=gb:D78132.1 gb:NM_005614.1 gb:AF148645.1	NM_005614		NP_005605

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201455_s_at	7.25E-04	Consensus includes gb:AJ132583.1 /DEF=Homo sapiens mRNA for puromycin sensitive aminopeptidase, partial. /FEA=mRNA /PROD=puromycin sensitive aminopeptidase /DB_XREF=gi:4210725 /UG=Hs.293007 aminopeptidase puromycin sensitive /FL=gb:NM_006310.1	NM_006310		NP_006301
201456_s_at	0.005119	BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)	AU160695	Hs.40323	NP_004716
201463_s_at	0.001406	gb:NM_006755.1 /DEF=Homo sapiens transaldolase 1 (TALDO1), mRNA. /FEA=mRNA /GEN=TALDO1 /PROD=transaldolase 1 /DB_XREF=gi:5803186 /UG=Hs.77290 transaldolase 1 /FL=gb:L19437.2 gb:NM_006755.1	NM_006755		NP_006746
201470_at	9.85E-04	gb:NM_004832.1 /DEF=Homo sapiens glutathione-S-transferase like; glutathione transferase omega (GSTTLp28), mRNA. /FEA=mRNA /GEN=GSTTLp28 /PROD=glutathione-S-transferase like /DB_XREF=gi:4758483 /UG=Hs.11465 glutathione-S-transferase like; glutathione transferase omega /FL=gb:BC000127.1 gb:U90313.1 gb:NM_004832.1 gb:AF212303.1	NM_004832		NP_004823

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201472_at	0.002559	gb:NM_003372.2 /DEF=Homo sapiens von Hippel-Lindau binding protein 1 (VBP1), mRNA. /FEA=mRNA /GEN=VBP1 /PROD=von Hippel-Lindau binding protein 1 /DB_XREF=gi:9257253 /UG=Hs.198307 von Hippel-Lindau binding protein 1 /FL=gb:U96759.1 gb:NM_003372.2	NM_003372		NP_003363
201475_x_at	0.01063	gb:NM_004990.1 /DEF=Homo sapiens methionine-tRNA synthetase (MARS), mRNA. /FEA=mRNA /GEN=MARS /PROD=methionine-tRNA synthetase /DB_XREF=gi:4826825 /UG=Hs.279946 methionine-tRNA synthetase /FL=gb:BC002384.1 gb:NM_004990.1 gb:D84224.1	NM_004990		NP_004981
201477_s_at	0.044397	gb:NM_001033.1 /DEF=Homo sapiens ribonucleotide reductase M1 polypeptide (RRM1), mRNA. /FEA=mRNA /GEN=RRM1 /PROD=ribonucleotide reductase M1 polypeptide /DB_XREF=gi:4506748 /UG=Hs.2934 ribonucleotide reductase M1 polypeptide /FL=gb:NM_001033.1	NM_001033		NP_001024
201482_at	0.006162	gb:NM_002826.2 /DEF=Homo sapiens quiescin Q6 (QSCN6), mRNA. /FEA=mRNA /GEN=QSCN6 /PROD=quiescin Q6 /DB_XREF=gi:13325074 /UG=Hs.77266 quiescin Q6 /FL=gb:L42379.1 gb:U97276.2 gb:NM_002826.2	NM_002826		NP_002817

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladd rCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201483_s_at	0.001812	gb:BC002802.1 /DEF=Homo sapiens, suppressor of Ty (S.cerevisiae) 4 homolog 1, clone MGC:3864, mRNA, complete cds. /FEA=mRNA /PROD=suppressor of Ty (S.cerevisiae) 4 homolog 1 /DB_XREF=gi:12803910 /UG=Hs.79058 suppressor of Ty (S.cerevisiae) 4 homolog 1 /FL=gb:BC002802.1 gb:U43923.1 gb:U38818.1 gb:U38817.1 gb:NM_003168.1	BC002802		NP_003159
201487_at	0.001406	gb:NM_001814.1 /DEF=Homo sapiens cathepsin C (CTSC), mRNA. /FEA=mRNA /GEN=CTSC /PROD=cathepsin C /DB_XREF=gi:4503140 /UG=Hs.10029 cathepsin C /FL=gb:NM_001814.1	NM_001814		NP_680475
201489_at	0.015807	gb:BC005020.1 /DEF=Homo sapiens, peptidylprolyl isomerase F (cyclophilin F), clone MGC:11022, mRNA, complete cds. /FEA=mRNA /PROD=peptidylprolyl isomerase F (cyclophilin F) /DB_XREF=gi:13477126 /UG=Hs.173125 peptidylprolyl isomerase F (cyclophilin F) /FL=gb:BC005020.1 gb:M80254.1 gb:NM_005729.1	BC005020		NP_005720
201491_at	0.007652	gb:NM_012111.1 /DEF=Homo sapiens chromosome 14 open reading frame 3 (C14ORF3), mRNA. /FEA=mRNA /GEN=C14ORF3 /PROD=chromosome 14 open reading frame 3 /DB_XREF=gi:6912279 /UG=Hs.204041 chromosome 14 open reading frame 3 /FL=gb:BC000321.1 gb:NM_012111.1 gb:AF164791.1	NM_012111		NP_036243
201493_s_at	0.001641	pumilio homolog 2 (Drosophila)	BE778078	Hs.6151	NP_056132

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201494_at	7.63E-04	gb:NM_005040.1 /DEF=Homo sapiens prolylcarboxypeptidase (angiotensinase C) (PRCP), mRNA. /FEA=mRNA /GEN=PRCP /PROD=prolylcarboxypeptidase (angiotensinase C) /DB_XREF=gi:4826939 /UG=Hs.75693 prolylcarboxypeptidase (angiotensinase C) /FL=gb:L13977.1 gb:NM_005040.1	NM_005040		NP_005031
201500_s_at	0.001942	gb:NM_021959.1 /DEF=Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 11 (PPP1R11), mRNA. /FEA=mRNA /GEN=PPP1R11 /PROD=protein phosphatase 1, regulatory (inhibitor) subunit 11 /DB_XREF=gi:11386174 /UG=Hs.82887 protein phosphatase 1, regulatory (inhibitor) subunit 11 /FL=gb:NM_021959.1	NM_021959		NP_740751
201507_at	0.04619	gb:NM_002622.2 /DEF=Homo sapiens prefoldin 1 (PFDN1), mRNA. /FEA=mRNA /GEN=PFDN1 /PROD=prefoldin 1 /DB_XREF=gi:12408673 /UG=Hs.132881 prefoldin 1 /FL=gb:NM_002622.2	NM_002622		NP_002613
201511_at	0.046228	gb:NM_001087.1 /DEF=Homo sapiens angio-associated, migratory cell protein (AAMP), mRNA. /FEA=mRNA /GEN=AAMP /PROD=angio-associated, migratory cell protein /DB_XREF=gi:4557228 /UG=Hs.83347 angio-associated, migratory cell protein /FL=gb:NM_001087.1 gb:M95627.1	NM_001087		NP_001078

Genes Corresponding To Differentially Expressed Genes in Figur 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201518_at	0.006722	gb:NM_006807.1 /DEF=Homo sapiens chromobox homolog 1 (Drosophila HP1 beta) (CBX1), mRNA. /FEA=mRNA /GEN=CBX1 /PROD=chromobox homolog 1 (Drosophila HP1 beta) /DB_XREF=gi:5803075 /UG=Hs.77254 chromobox homolog 1 (Drosophila HP1 beta) /FL=gb:U35451.1 gb:BC002609.1 gb:NM_006807.1	NM_006807		NP_006798
201527_at	0.007271	gb:NM_004231.1 /DEF=Homo sapiens ATPase, vacuolar, 14 kD (ATP6S14), mRNA. /FEA=mRNA /GEN=ATP6S14 /PROD=ATPase, vacuolar, 14 kD /DB_XREF=gi:4757819 /UG=Hs.78089 ATPase, vacuolar, 14 kD /FL=gb:D49400.1 gb:NM_004231.1	NM_004231		NP_004222
201533_at	0.034296	gb:NM_001904.1 /DEF=Homo sapiens catenin (cadherin-associated protein), beta 1 (88kD) (CTNNB1), mRNA. /FEA=mRNA /GEN=CTNNB1 /PROD=catenin (cadherin-associated protein), beta 1(88kD) /DB_XREF=gi:4503130 /UG=Hs.171271 catenin (cadherin-associated protein), beta 1 (88kD) /FL=gb:NM_001904.1	NM_001904		NP_001895
201535_at	0.005652	gb:NM_007106.1 /DEF=Homo sapiens ubiquitin-like 3 (UBL3), mRNA. /FEA=mRNA /GEN=UBL3 /PROD=ubiquitin-like 3 /DB_XREF=gi:6005927 /UG=Hs.173091 ubiquitin-like 3 /FL=gb:AF044221.1 gb:AL080177.1 gb:NM_007106.1	NM_007106		NP_009037

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gen Accession No.	Unigen Accession No.	Prot in Accession No.
201541_s_at	0.005189	gb:NM_006349.1 /DEF=Homo sapiens putative cyclin G1 interacting protein. (CG1I), mRNA. /FEA=mRNA /GEN=CG1I /PROD=putative cyclin G1 interacting protein /DB_XREF=gi:5453616 /UG=Hs.10028 putative cyclin G1 interacting protein /FL=gb:U61837.1 gb:NM_006349.1	NM_006349		NP_006340
201543_s_at	0.040891	gb:NM_020150.1 /DEF=Homo sapiens SAR1 protein (SAR1), mRNA. /FEA=mRNA /GEN=SAR1 /PROD=SAR1 protein /DB_XREF=gi:9910541 /UG=Hs.110796 SAR1 protein /FL=gb:AY008268.1 gb:AL136724.1 gb:AF261717.1 gb:NM_020150.1	NM_020150		
201546_at	6.66E-04	gb:NM_004238.1 /DEF=Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA. /FEA=mRNA /GEN=TRIP12 /PROD=thyroid hormone receptor interactor 12 /DB_XREF=gi:10863902 /UG=Hs.138617 thyroid hormone receptor interactor 12 /FL=gb:NM_004238.1 gb:D28476.1	NM_004238		NP_004229
201549_x_at	0.006864	gb:NM_006618.1 /DEF=Homo sapiens putative DNAchromatin binding motif (PLU-1), mRNA. /FEA=mRNA /GEN=PLU-1 /PROD=putative DNAchromatin binding motif /DB_XREF=gi:5729977 /UG=Hs.143323 putative DNAchromatin binding motif /FL=gb:NM_006618.1	NM_006618		NP_006609

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
201552_at	7.63E-04	gb:NM_005561.2 /DEF=Homo sapiens lysosomal-associated membrane protein 1 (LAMP1), mRNA. /FEA=mRNA /GEN=LAMP1 /PROD=lysosomal-associated membrane protein 1 /DB_XREF=gi:7669500 /UG=Hs.150101 lysosomal-associated membrane protein 1 /FL=gb:J04182.1 gb:J03263.1 gb:NM_005561.2	NM_005561		NP_005552
201553_s_at	0.008506	gb:NM_005561.2 /DEF=Homo sapiens lysosomal-associated membrane protein 1 (LAMP1), mRNA. /FEA=mRNA /GEN=LAMP1 /PROD=lysosomal-associated membrane protein 1 /DB_XREF=gi:7669500 /UG=Hs.150101 lysosomal-associated membrane protein 1 /FL=gb:J04182.1 gb:J03263.1 gb:NM_005561.2	NM_005561		NP_005552
201554_x_at	7.63E-04	gb:NM_004130.1 /DEF=Homo sapiens glycogenin (GYG), mRNA. /FEA=mRNA /GEN=GYG /PROD=glycogenin /DB_XREF=gi:4758491 /UG=Hs.174071 glycogenin /FL=gb:U44131.1 gb:BC000033.1 gb:NM_004130.1 gb:U31525.1	NM_004130		NP_004121
201558_at	7.63E-04	gb:NM_003610.1 /DEF=Homo sapiens RAE1 (RNA export 1, S.pombe) homolog (RAE1), mRNA. /FEA=mRNA /GEN=RAE1 /PROD=RAE1 (RNA export 1, S.pombe) homolog /DB_XREF=gi:4506398 /UG=Hs.196209 RAE1 (RNA export 1, S.pombe) homolog /FL=gb:U84720.1 gb:NM_003610.1	NM_003610		NP_003601

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
201561_s_at	0.010937	gb:NM_014944.1 /DEF=Homo sapiens KIAA0911 protein (KIAA0911), mRNA. /FEA=mRNA /GEN=KIAA0911 /PROD=KIAA0911 protein /DB_XREF=gi:7662373 /UG=Hs.29665 KIAA0911 protein /FL=gb:AB020718.1 gb:NM_014944.1	NM_014944		NP_055759
201563_at	0.029279	gb:L29008.1 /DEF=Human L-iditol-2 dehydrogenase mRNA, complete cds. /FEA=mRNA /PROD=L-iditol-2 dehydrogenase /DB_XREF=gi:496077 /UG=Hs.878 sorbitol dehydrogenase /FL=gb:NM_003104.1 gb:L29008.1 gb:U07361.1	L29008		NP_003095
201565_s_at	0.017199	gb:NM_002166.1 /DEF=Homo sapiens inhibitor of DNA binding 2, dominant negative helix-loop-helix protein (ID2), mRNA. /FEA=mRNA /GEN=ID2 /PROD=inhibitor of DNA binding 2, dominant negative helix-loop-helix protein /DB_XREF=gi:4504570 /UG=Hs.180919 inhibitor of DNA binding 2, dominant negative helix-loop-helix protein /FL=gb:M97796.1 gb:NM_002166.1 gb:D13891.1	NM_002166		NP_002157
201566_x_at	0.02937	gb:D13891.1 /DEF=Human mRNA for Id-2H, complete cds. /FEA=mRNA /GEN=Id-2H /PROD=Id-2H /DB_XREF=gi:464183 /UG=Hs.180919 inhibitor of DNA binding 2, dominant negative helix-loop-helix protein /FL=gb:M97796.1 gb:NM_002166.1 gb:D13891.1	D13891		NP_002157

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
201573_s_at	0.001181	gb:M75715.1 /DEF=Human TB3-1 mRNA, complete cds. /FEA=mRNA /PROD=TB3-1 /DB_XREF=gi:338686 /UG=Hs.77324 eukaryotic translation termination factor 1 /FL=gb:U90176.1 gb:M75715.1 gb:NM_004730.1	M75715		NP_004721
201574_at	0.006596	gb:NM_004730.1 /DEF=Homo sapiens eukaryotic translation termination factor 1 (ETF1), mRNA. /FEA=mRNA /GEN=ETF1 /PROD=eukaryotic translation termination factor 1 /DB_XREF=gi:4759033 /UG=Hs.77324 eukaryotic translation termination factor 1 /FL=gb:U90176.1 gb:M75715.1 gb:NM_004730.1	NM_004730		NP_004721
201575_at	0.045986	gb:NM_012245.1 /DEF=Homo sapiens SKI-INTERACTING PROTEIN (SNW1), mRNA. /FEA=mRNA /GEN=SNW1 /PROD=SKI-INTERACTING PROTEIN /DB_XREF=gi:6912675 /UG=Hs.79008 SKI-INTERACTING PROTEIN /FL=gb:U51432.1 gb:AF045184.1 gb:NM_012245.1	NM_012245		NP_036377
201576_s_at	6.66E-04	gb:NM_000404.1 /DEF=Homo sapiens galactosidase, beta 1 (GLB1), mRNA. /FEA=mRNA /GEN=GLB1 /PROD=galactosidase, beta 1 /DB_XREF=gi:10834965 /UG=Hs.79222 galactosidase, beta 1 /FL=gb:NM_000404.1 gb:M27507.1 gb:M22590.1 gb:M34423.1	NM_000404		NP_000395
201581_at	0.018248	hypothetical protein DJ971N18.2	AL544094	Hs.169358	NP_066979

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201582_at	0.005758	Consensus includes gb:AL121900 /DEF=Human DNA sequence from clone RP11-379J5 on chromosome 20 Contains the last exon of the SEC23B gene for Sec23 (S. cerevisiae) homolog B, a putative novel gene, the 5' end of the gene for a novel protein similar to bacterial histidyl-tRNA synthetase... /FEA=mRNA /DB_XREF=gi:11121203 /UG=Hs.173497 Sec23 (S. cerevisiae) homolog B /FL=gb:BC005404.1 gb:NM_006363.1	AL121900		
201583_s_at	7.63E-04	gb:NM_006363.1 /DEF=Homo sapiens Sec23 (S. cerevisiae) homolog B (SEC23B), mRNA. /FEA=mRNA /GEN=SEC23B /PROD=Sec23 (S. cerevisiae) homolog B /DB_XREF=gi:5454043 /UG=Hs.173497 Sec23 (S. cerevisiae) homolog B /FL=gb:BC005404.1 gb:NM_006363.1	NM_006363		NP_116781
201584_s_at	0.049392	gb:NM_005804.1 /DEF=Homo sapiens nuclear RNA helicase, DECD variant of DEAD box family (DDXL), mRNA. /FEA=mRNA /GEN=DDXL /PROD=nuclear RNA helicase, DECD variant of DEAD boxfamily /DB_XREF=gi:5031658 /UG=Hs.179606 nuclear RNA helicase, DECD variant of DEAD box family /FL=gb:BC001009.1 gb:U90426.1 gb:NM_005804.1	NM_005804		NP_620551
201585_s_at	0.049392	splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated)	BG035151	Hs.180610	NP_005057

G n s Corresponding To Differentially Expr ssed G n s in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201586_s_at	0.003393	gb:NM_005066.1 /DEF=Homo sapiens splicing factor prolineglutamine rich (polypyrimidine tract-binding protein-associated) (SFPQ), mRNA. /FEA=mRNA /GEN=SFPQ /PROD=splicing factor prolineglutamine rich(polypyrimidine tract-binding protein-associated) /DB_XREF=gi:4826997 /UG=Hs.180610 splicing factor prolineglutamine rich (polypyrimidine tract-binding protein-associated) /FL=gb:NM_005066.1	NM_005066		NP_005057
201587_s_at	0.029624	gb:NM_001569.2 /DEF=Homo sapiens interleukin-1 receptor-associated kinase 1 (IRAK1), mRNA. /FEA=mRNA /GEN=IRAK1 /PROD=interleukin-1 receptor-associated kinase 1 /DB_XREF=gi:4755143 /UG=Hs.182018 interleukin-1 receptor-associated kinase 1 /FL=gb:L76191.1 gb:NM_001569.2	NM_001569		NP_001560
201588_at	0.007591	gb:NM_004786.1 /DEF=Homo sapiens thioredoxin-like, 32kD (TXNL), mRNA. /FEA=mRNA /GEN=TXNL /PROD=thioredoxin-like, 32kD /DB_XREF=gi:4759273 /UG=Hs.18792 thioredoxin-like, 32kD /FL=gb:BC001156.1 gb:AF003938.1 gb:AF051896.1 gb:AF052659.1 gb:NM_004786.1	NM_004786		NP_004777

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201589_at	0.002749	Consensus includes gb:D80000.1 /DEF=Human mRNA for KIAA0178 gene, partial cds. /FEA=mRNA /GEN=KIAA0178 /DB_XREF=gi:1136415 /UG=Hs.211602.SMC1 (structural maintenance of chromosomes 1, yeast)-like 1 /FL=gb:NM_006306.1	D80000		NP_006297
201593_s_at	0.001835	uncharacterized hypothalamus protein HT010	AV716798	Hs.6375	NP_060941
201594_s_at	0.005119	gb:NM_005134.1 /DEF=Homo sapiens protein phosphatase 4, regulatory subunit 1 (PPP4R1), mRNA. /FEA=mRNA /GEN=PPP4R1 /PROD=protein phosphatase 4, regulatory subunit 1 /DB_XREF=gi:4826933 /UG=Hs.3382 protein phosphatase 4, regulatory subunit 1 /FL=gb:AF111106.1 gb:NM_005134.1 gb:AF100744.1	NM_005134		NP_005125
201597_at	0.030968	gb:NM_001865.1 /DEF=Homo sapiens cytochrome c oxidase subunit VIIa polypeptide 2 (liver) (COX7A2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=COX7A2 /PROD=cytochrome c oxidase subunit VIIa polypeptide 2(liver) /DB_XREF=gi:4502988 /UG=Hs.70312 cytochrome c oxidase subunit VIIa polypeptide 2 (liver) /FL=gb:NM_001865.1	NM_001865		NP_001856

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201599_at	7.63E-04	gb:NM_000274.1 /DEF=Homo sapiens ornithine aminotransferase (gyrate atrophy) (OAT), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=OAT /PROD=ornithine aminotransferase precursor /DB_XREF=gi:4557808 /UG=Hs.75485 ornithine aminotransferase (gyrate atrophy) /FL=gb:BC000964.1 gb:M12267.1 gb:M23204.1 gb:M14963.1 gb:NM_000274.1	NM_000274		NP_000265
201602_s_at	0.003315	Consensus includes gb:BE737620 /FEA=EST /DB_XREF=gi:10151612 /DB_XREF=est:601572895F1 /CLONE=IMAGE:3839831 /UG=Hs.16533 myosin phosphatase, target subunit 1 /FL=gb:NM_002480.1	NM_002480		NP_002471
201603_at	0.003965	Consensus includes gb:AI817061 /FEA=EST /DB_XREF=gi:5436140 /DB_XREF=est:wj76e05.x1 /CLONE=IMAGE:2408768 /UG=Hs.16533 myosin phosphatase, target subunit 1 /FL=gb:NM_002480.1	NM_002480		NP_002471
201604_s_at	0.038202	gb:NM_002480.1 /DEF=Homo sapiens myosin phosphatase, target subunit 1 (MYPT1), mRNA. /FEA=mRNA /GEN=MYPT1 /PROD=myosin phosphatase target subunit 1 /DB_XREF=gi:4505316 /UG=Hs.16533 myosin phosphatase, target subunit 1 /FL=gb:NM_002480.1	NM_002480		NP_002471
201606_s_at	0.005652	nuclear phosphoprotein similar to <i>S. cerevisiae</i> PWP1	BE796924	Hs.172589	NP_008993
201609_x_at	0.012369	isoprenylcysteine carboxyl methyltransferase	AL578502	Hs.183212	NP_733806

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201619_at	7.63E-04	gb:NM_006793.1 /DEF=Homo sapiens peroxiredoxin 3 (PRDX3), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=PRDX3 /PROD=peroxiredoxin 3 /DB_XREF=gi:5802973 /UG=Hs.75454 peroxiredoxin 3 /FL=gb:BC002685.1 gb:NM_006793.1 gb:D49396.1	NM_006793		NP_006784
201623_s_at	0.010277	gb:BC000629.1 /DEF=Homo sapiens, Similar to aspartyl-tRNA synthetase, clone MGC:1562, mRNA, complete cds. /FEA=mRNA /PROD=Similar to aspartyl-tRNA synthetase /DB_XREF=gi:12653688 /UG=Hs.80758 aspartyl-tRNA synthetase /FL=gb:BC000629.1 gb:J05032.1 gb:NM_001349.1	BC000629		NP_001340
201625_s_at	0.022311	insulin induced gene 1	BE300521	Hs.56205	NP_005533
201626_at	0.01747	insulin induced gene 1	BE300521	Hs.56205	NP_005533
201627_s_at	7.76E-04	gb:NM_005542.1 /DEF=Homo sapiens insulin induced gene 1 (INSIG1), mRNA. /FEA=mRNA /GEN=INSIG1 /PROD=insulin induced gene 1 /DB_XREF=gi:5031800 /UG=Hs.56205 insulin induced gene 1 /FL=gb:NM_005542.1	NM_005542		NP_005533
201628_s_at	6.66E-04	gb:NM_006570.1 /DEF=Homo sapiens Ras-related GTP-binding protein (RAGA), mRNA. /FEA=mRNA /GEN=RAGA /PROD=Ras-related GTP-binding protein /DB_XREF=gi:5729998 /UG=Hs.57304 Ras-related GTP-binding protein /FL=gb:U41654.1 gb:NM_006570.1	NM_006570		NP_006561

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201631_s_at	0.024403	gb:NM_003897.1 /DEF=Homo sapiens immediate early response 3 (IER3), mRNA. /FEA=mRNA /GEN=IER3 /PROD=immediate early response 3 /DB_XREF=gi:4503328 /UG=Hs.76095 immediate early response 3 /FL=gb:BC000844.1 gb:BC005080.1 gb:AF083421.1 gb:NM_003897.1	NM_003897		NP_434702
201642_at	0.001852	gb:NM_005534.1 /DEF=Homo sapiens interferon gamma receptor 2 (interferon gamma transducer 1) (IFNGR2), mRNA. /FEA=mRNA /GEN=IFNGR2 /PROD=interferon gamma receptor 2 (interferon gamma transducer 1) /DB_XREF=gi:5031782 /UG=Hs.177559 interferon gamma receptor 2 (interferon gamma transducer 1) /FL=gb:BC003624.1 gb:U05875.1 gb:U05877.1 gb:NM_005534.1	NM_005534		NP_005525
201643_x_at	0.001406	gb:NM_016604.1 /DEF=Homo sapiens putative zinc finger protein (LOC51780), mRNA. /FEA=mRNA /GEN=LOC51780 /PROD=putative zinc finger protein /DB_XREF=gi:7706598 /UG=Hs.24125 putative zinc finger protein /FL=gb:AF251039.1 gb:NM_016604.1	NM_016604		NP_057688
201648_at	0.007503	Janus kinase 1 (a protein tyrosine kinase)	AL039831	Hs.50651	

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201651_s_at	0.002603	gb:NM_007229.1 /DEF=Homo sapiens protein kinase C and casein kinase substrate in neurons 2 (PACIN2), mRNA. /FEA=mRNA /GEN=PACIN2 /PROD=protein kinase C and casein kinase substrate in neurons 2 /DB_XREF=gi:6005825 /UG=Hs.18842 protein kinase C and casein kinase substrate in neurons 2 /FL=gb:AF128536.1 gb:NM_007229.1	NM_007229		NP_009160
201652_at	0.002692	gb:NM_006837.1 /DEF=Homo sapiens COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 5 (COPS5), mRNA. /FEA=mRNA /GEN=COPS5 /PROD=COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 5 /DB_XREF=gi:5803045 /UG=Hs.198767 COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 5 /FL=gb:BC001187.1 gb:BC001859.1 gb:U65928.1 gb:U70734.1 gb:NM_006837.1	NM_006837		NP_006828
201662_s_at	0.003976	gb:D89053.1 /DEF=Homo sapiens mRNA for Acyl-CoA synthetase 3, complete cds. /FEA=mRNA /PROD=Acyl-CoA synthetase 3 /DB_XREF=gi:4165017 /UG=Hs.268012 fatty-acid-Coenzyme A ligase, long-chain 3 /FL=gb:NM_004457.2 gb:D89053.1 gb:AF116690.1	D89053		NP_004448

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201666_at	6.66E-04	gb:NM_003254.1 /DEF=Homo sapiens tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor) (TIMP1), mRNA. /FEA=mRNA /GEN=TIMP1 /PROD=tissue inhibitor of metalloproteinase 1 precursor /DB_XREF=gi:4507508 /UG=Hs.5831 tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor) /FL=gb:BC000866.1 gb:M12670.1 gb:M59906.1 gb:NM_003254.1	NM_003254		NP_003245
201669_s_at	6.66E-04	gb:NM_002356.4 /DEF=Homo sapiens myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L) (MACS), mRNA. /FEA=mRNA /GEN=MACS /PROD=myristoylated alanine-rich protein kinase C substrate /DB_XREF=gi:11125771 /UG=Hs.75607 myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L) /FL=gb:NM_002356.4 gb:M68956.1 gb:D10522.1	NM_002356		NP_002347
201670_s_at	0.001719	gb:M68956.1 /DEF=Human myristoylated alanine-rich C-kinase substrate mRNA, complete cds. /FEA=mRNA /GEN=MACS /PROD=myristoylated alanine-rich C-kinase substrate /DB_XREF=gi:187386 /UG=Hs.75607 myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L) /FL=gb:NM_002356.4 gb:M68956.1 gb:D10522.1	M68956		NP_002347

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201671_x_at	0.040119	gb:BC003556.1 /DEF=Homo sapiens, ubiquitin specific protease 14 (tRNA-guanine transglycosylase), clone MGC:1453, mRNA, complete cds. /FEA=mRNA) /PROD=ubiquitin specific protease 14 (tRNA-guanine transglycosylase) /DB_XREF=gi:13097695 /UG=Hs.75981 ubiquitin specific protease 14 (tRNA-guanine transglycosylase) /FL=gb:BC003556.1 gb:NM_005151.1 gb:U30888.1	BC003556		NP_005142
201673_s_at	0.048089	gb:NM_002103.1 /DEF=Homo sapiens glycogen synthase 1 (muscle) (GYS1), mRNA. /FEA=mRNA /GEN=GYS1 /PROD=glycogen synthase 1 (muscle) /DB_XREF=gi:4504232 /UG=Hs.772 glycogen synthase 1 (muscle) /FL=gb:U32573.1 gb:BC002617.1 gb:J04501.1 gb:NM_002103.1	NM_002103		NP_002094
201684_s_at	0.001942	KIAA0737 gene product	BE783632	Hs.194035	NP_055643
201688_s_at	0.027454	tumor protein D52	BE974098	Hs.2384	NP_005070
201690_s_at	0.038384	tumor protein D52	BE974098	Hs.2384	NP_005070
201695_s_at	9.85E-04	gb:NM_000270.1 /DEF=Homo sapiens nucleoside phosphorylase (NP), mRNA. /FEA=mRNA /GEN=NP /PROD=purine nucleoside phosphorylase /DB_XREF=gi:4557800 /UG=Hs.75514 nucleoside phosphorylase /FL=gb:NM_000270.1	NM_000270		NP_000261

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201698_s_at	7.63E-04	gb:NM_003769.1 /DEF=Homo sapiens splicing factor, arginineserine-rich 9 (SFRS9), mRNA. /FEA=mRNA /GEN=SFRS9 /PROD=splicing factor, arginineserine-rich 9 /DB_XREF=gi:4506902 /UG=Hs.77608 splicing factor, arginineserine-rich 9 /FL=gb:U30825.1 gb:NM_003769.1	NM_003769		NP_003760
201699_at	0.014372	gb:NM_002806.1 /DEF=Homo sapiens' proteasome (prosome, macropain) 26S subunit, ATPase, 6 (PSMC6), mRNA. /FEA=mRNA /GEN=PSMC6 /PROD=proteasome (prosome, macropain) 26S subunit, ATPase, 6 /DB_XREF=gi:4506214 /UG=Hs.79357 proteasome (prosome, macropain) 26S subunit, ATPase, 6 /FL=gb:BC005390.1 gb:D78275.1 gb:AF006305.1 gb:NM_002806.1	NM_002806		NP_002797
201700_at	0.041179	gb:NM_001760.1 /DEF=Homo sapiens cyclin D3 (CCND3), mRNA. /FEA=mRNA /GEN=CCND3 /PROD=cyclin D3 /DB_XREF=gi:4502618 /UG=Hs.83173 cyclin D3 /FL=gb:M90814.1 gb:M92287.1 gb:NM_001760.1	NM_001760		NP_001751

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201703_s_at	0.003366	gb:NM_002714.1 /DEF=Homo sapiens protein phosphatase 1, regulatory subunit 10 (PPP1R10), mRNA. /FEA=mRNA /GEN=PPP1R10 /PROD=protein phosphatase 1, regulatory subunit 10 /DB_XREF=gi:4506008 /UG=Hs.106019 protein phosphatase 1, regulatory subunit 10 /FL=gb:NM_002714.1	NM_002714		NP_002705
201705_at	0.005574	gb:NM_002811.1 /DEF=Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 7 (Mov34 homolog) (PSMD7), mRNA. /FEA=mRNA /GEN=PSMD7 /PROD=proteasome (prosome, macropain) 26S subunit, non-ATPase, 7 (Mov34 homolog) /DB_XREF=gi:4506230 /UG=Hs.155543 proteasome (prosome, macropain) 26S subunit, non-ATPase, 7 (Mov34 homolog) /FL=gb:NM_002811.1 gb:D50063.1	NM_002811		NP_002802
201713_s_at	0.009678	gb:D42063.1 /DEF=Human mRNA for RanBP2 (Ran-binding protein 2), complete cds. /FEA=mRNA /PROD=RanBP2 (Ran-binding protein 2) /DB_XREF=gi:924266 /UG=Hs.199179 RAN binding protein 2 /FL=gb:NM_006267.2 gb:D42063.1	D42063		NP_006258

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
201716_at	6.66E-04	gb:NM_003099.1 /DEF=Homo sapiens sorting nexin 1 (SNX1), mRNA. /FEA=mRNA /GEN=SNX1 /PROD=sorting nexin 1 /DB_XREF=gi:4507138 /UG=Hs.75283 sorting nexin 1 /FL=gb:BC000357.1 gb:U53225.1 gb:AF065483.1 gb:NM_003099.1	NM_003099		NP_690039
201722_s_at	6.66E-04	Consensus includes gb:AV692127 /FEA=EST /DB_XREF=gi:10293990 /DB_XREF=est:AV692127 /CLONE=GKCAOB04 /UG=Hs.80120 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1) /FL=gb:U41514.1 gb:NM_020474.2	NM_020474		NP_065207
201724_s_at	0.00158	gb:NM_020474.2 /DEF=Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1) (GALNT1), mRNA. /FEA=mRNA /GEN=GALNT1 /PROD=polypeptide N-acetylgalactosaminyltransferase 1 /DB_XREF=gi:13124890 /UG=Hs.80120 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1) /FL=gb:U41514.1 gb:NM_020474.2	NM_020474		NP_065207

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
201725_at	0.00158	gb:NM_006023.1 /DEF=Homo sapiens D123 gene product (D123), mRNA. /FEA=mRNA /GEN=D123 /PROD=D123 gene product /DB_XREF=gi:5174422 /UG=Hs.82043 D123 gene product /FL=gb:BC001600.1 gb:D14878.1 gb:U27112.1 gb:NM_006023.1	NM_006023		NP_006014
201730_s_at	0.029624	translocated promoter region (to activated MET oncogene)	BF110993	Hs.169750	NP_003283
201737_s_at	0.049282	gb:NM_005885.1 /DEF=Homo sapiens similar to S. cerevisiae SSM4 (TEB4), mRNA. /FEA=mRNA /GEN=TEB4 /PROD=similar to S. cerevisiae SSM4 /DB_XREF=gi:5032166 /UG=Hs.20141 similar to S. cerevisiae SSM4 /FL=gb:AF009301.1 gb:NM_005885.1	NM_005885		NP_005876
201738_at	6.66E-04	gb:NM_005875.1 /DEF=Homo sapiens translation factor. sui1 homolog (GC20), mRNA. /FEA=mRNA /GEN=GC20 /PROD=translation factor sui1 homolog /DB_XREF=gi:5031710 /UG=Hs.21756 translation factor sui1 homolog /FL=gb:AF064607.1 gb:AF077052.1 gb:NM_005875.1 gb:AF263452.1	NM_005875		NP_005866
201745_at	0.007895	gb:NM_002822.1 /DEF=Homo sapiens protein tyrosine kinase 9 (PTK9), mRNA. /FEA=mRNA /GEN=PTK9 /PROD=protein tyrosine kinase 9 /DB_XREF=gi:4506274 /UG=Hs.82643 protein tyrosine kinase 9 /FL=gb:NM_002822.1 gb:U02680.1	NM_002822		NP_002813

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201758_at	6.66E-04	gb:NM_006292.1 /DEF=Homo sapiens tumor susceptibility gene 101 (TSG101), mRNA. /FEA=mRNA /GEN=TSG101 /PROD=tumor susceptibility gene 101 /DB_XREF=gi:5454139 /UG=Hs.118910 tumor susceptibility gene 101 /FL=gb:BC002487.1 gb:U82130.1 gb:NM_006292.1	NM_006292		NP_006283
201760_s_at	6.66E-04	gb:NM_018639.1 /DEF=Homo sapiens CS box-containing WD protein (LOC55884), mRNA. /FEA=mRNA /GEN=LOC55884 /PROD=CS box-containing WD protein /DB_XREF=gi:8923880 /UG=Hs.136644 CS box-containing WD protein /FL=gb:AF229181.1 gb:AF163324.1 gb:NM_018639.1	NM_018639		NP_061109
201761_at	0.003373	gb:NM_006636.2 /DEF=Homo sapiens methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase (MTHFD2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=MTHFD2 /PROD=methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase precursor /DB_XREF=gi:13699869 /UG=Hs.154672 methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase /FL=gb:NM_006636.2	NM_006636		NP_006627
201765_s_at	0.023856	hexosaminidase A (alpha polypeptide)	AL523158	Hs.119403	NP_000511

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201769_at	7.63E-04	gb:NM_014666.1 /DEF=Homo sapiens KIAA0171 gene product (KIAA0171), mRNA. /FEA=mRNA /GEN=KIAA0171 /PROD=KIAA0171 gene product /DB_XREF=gi:7661967 /UG=Hs.155623 KIAA0171 gene product /FL=gb:D79993.1 gb:BC004467.1 gb:NM_014666.1	NM_014666		NP_055481
201772_at	0.003842	gb:NM_015878.1 /DEF=Homo sapiens antizyme inhibitor (LOC51582), mRNA. /FEA=mRNA /GEN=LOC51582 /PROD=antizyme inhibitor. /DB_XREF=gi:7706219 /UG=Hs.223014 antizyme inhibitor /FL=gb:D88674.1 gb:NM_015878.1	NM_015878		NP_680479
201773_at	0.003305	gb:NM_015339.1 /DEF=Homo sapiens activity-dependent neuroprotective protein (ADNP), mRNA. /FEA=mRNA /GEN=ADNP /PROD=activity-dependent neuroprotective protein. /DB_XREF=gi:12229216 /UG=Hs.3657 activity-dependent neuroprotective protein /FL=gb:AF250860.1 gb:NM_015339.1	NM_015339		NP_056154
201774_s_at	0.048144	Consensus includes gb:AK022511.1 /DEF=Homo sapiens cDNA FLJ12449 fis, clone NT2RM1000314, highly similar to Human mRNA for KIAA0159 gene. /FEA=mRNA /DB_XREF=gi:10433940 /UG=Hs.5719 chromosome condensation-related SMC-associated protein 1 /FL=gb:NM_014865.1 gb:D63880.1	AK022511		NP_055680

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201776_s_at	0.015926	Consensus includes gb:AK001487.1 /DEF=Homo sapiens cDNA FLJ10625 fis, clone NT2RP2005540, highly similar to Homo sapiens mRNA for KIAA0494 protein. /FEA=mRNA /DB_XREF=gi:7022773 /UG=Hs.62515 KIAA0494 gene product /FL=gb:BC002525.1 gb:AB007963.1 gb:NM_014774.1	AK001487		NP_055589
201780_s_at	7.25E-04	gb:NM_007282.1 /DEF=Homo sapiens ring finger protein 13 (RNF13), mRNA. /FEA=mRNA /GEN=RNF13 /PROD=ring finger protein 13 /DB_XREF=gi:6005863 /UG=Hs.6900 ring finger protein 13 /FL=gb:AF037204.1 gb:AF070558.1 gb:NM_007282.1	NM_007282		NP_009213
201784_s_at	0.013527	gb:NM_014267.1 /DEF=Homo sapiens small acidic protein (IMAGE145052), mRNA. /FEA=mRNA /GEN=IMAGE145052 /PROD=small acidic protein /DB_XREF=gi:7657233 /UG=Hs.78050 small acidic protein /FL=gb:U51678.1 gb:NM_014267.1	NM_014267		NP_055082
201794_s_at	0.034006	gb:NM_014837.1 /DEF=Homo sapiens KIAA0250 gene product (KIAA0250), mRNA. /FEA=mRNA /GEN=KIAA0250 /PROD=KIAA0250 gene product /DB_XREF=gi:7662023 /UG=Hs.15087 KIAA0250 gene product /FL=gb:D87437.1 gb:NM_014837.1	NM_014837		NP_055652

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201795_at	0.018901	gb:NM_002296.1 /DEF=Homo sapiens lamin B receptor (LBR), mRNA. /FEA=mRNA /GEN=LBR /PROD=lamin B receptor /DB_XREF=gi:4504960 /UG=Hs.152931 lamin B receptor /FL=gb:L25931.1 gb:NM_002296.1	NM_002296		NP_002287
201805_at	0.01263	gb:NM_002733.1 /DEF=Homo sapiens protein kinase, AMP-activated, gamma 1 non-catalytic subunit (PRKAG1), mRNA. /FEA=mRNA /GEN=PRKAG1 /PROD=protein kinase, AMP-activated, gamma 1 non-catalytic subunit /DB_XREF=gi:4506060 /UG=Hs.3136 protein kinase, AMP-activated, gamma 1 non-catalytic subunit /FL=gb:BC000358.1 gb:U42412.1 gb:NM_002733.1	NM_002733		NP_002724
201807_at	0.001641	gb:NM_004896.1 /DEF=Homo sapiens vacuolar protein sorting 26 (yeast homolog) (VPS26), mRNA. /FEA=mRNA /GEN=VPS26 /PROD=vacuolar protein sorting 26 (yeast homolog) /DB_XREF=gi:4758509 /UG=Hs.67052 vacuolar protein sorting 26 (yeast homolog) /FL=gb:AF054179.1 gb:NM_004896.1 gb:AF175266.1	NM_004896		NP_004887

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201811_x_at	9.85E-04	gb:NM_004844.1 /DEF=Homo sapiens SH3-domain binding protein 5 (BTK-associated) (SH3BP5), mRNA. /FEA=mRNA /GEN=SH3BP5 /PROD=SH3-domain binding protein 5 (BTK-associated) /DB_XREF=gi:4759057 /UG=Hs.109150 SH3-domain binding protein 5 (BTK-associated) /FL=gb:AB005047.1 gb:NM_004844.1	NM_004844		NP_004835
201814_at	0.027219	Consensus includes gb:AI300084 /FEA=EST /DB_XREF=gi:3959430 /DB_XREF=est:qn59f02.x1 /CLONE=IMAGE:1902555 /UG=Hs.115740 KIAA0210 gene product /FL=gb:D86965.1 gb:NM_014744.1	NM_014744		NP_055559
201815_s_at	0.044483	gb:NM_014744.1 /DEF=Homo sapiens KIAA0210 gene product (KIAA0210), mRNA. /FEA=mRNA /GEN=KIAA0210 /PROD=KIAA0210 gene product /DB_XREF=gi:7661997 /UG=Hs.115740 KIAA0210 gene product /FL=gb:D86965.1 gb:NM_014744.1	NM_014744		NP_055559
201816_s_at	0.005347	gb:NM_001483.1 /DEF=Homo sapiens glioblastoma amplified sequence (GBAS), mRNA. /FEA=mRNA /GEN=GBAS /PROD=glioblastoma amplified sequence /DB_XREF=gi:4503936 /UG=Hs.152707 glioblastoma amplified sequence /FL=gb:BC000732.1 gb:BC001837.1 gb:AF029786.1 gb:NM_001483.1	NM_001483		NP_001474

Genes Corresponding To Differentially Expressed Genes in Figur 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201817_at	0.001763	gb:NM_014671.1 /DEF=Homo sapiens KIAA0010 gene product (KIAA0010), mRNA. /FEA=mRNA /GEN=KIAA0010 /PROD=KIAA0010 gene product /DB_XREF=gi:7661855 /UG=Hs.155287 KIAA0010 gene product /FL=gb:D13635.1 gb:NM_014671.1	NM_014671		NP_055486
201821_s_at	0.026556	gb:BC004439.1 /DEF=Homo sapiens, translocase of inner mitochondrial membrane 17 (yeast) homolog A, clone MGC:4031, mRNA, complete cds. /FEA=mRNA /PROD=translocase of inner mitochondrial membrane 17(yeast) homolog A /DB_XREF=gi:13325239 /UG=Hs.20716 translocase of inner mitochondrial membrane 17 (yeast) homolog A /FL=gb:BC004439.1 gb:AF106622.1 gb:NM_006335.1	BC004439		NP_006326
201823_s_at	0.004519	gb:NM_004290.1 /DEF=Homo sapiens ring finger protein 14 (RNF14), mRNA. /FEA=mRNA /GEN=RNF14 /PROD=ring finger protein 14 /DB_XREF=gi:4757761 /UG=Hs.215857 ring finger protein 14 /FL=gb:AF060544.1 gb:NM_004290.1 gb:AB022663.1	NM_004290		NP_004281

Genes Corresponding To Differentially Expressed Genes in Figur 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201824_at	0.03867	gb:AB022663.1 /DEF=Homo sapiens HFB30 mRNA, complete cds. /FEA=mRNA /GEN=HFB30 /DB_XREF=gi:5019617 /UG=Hs.215857 ring finger protein 14 /FL=gb:AF060544.1 gb:NM_004290.1 gb:AB022663.1	AB022663		NP_004281
201826_s_at	0.020839	gb:NM_016002.1 /DEF=Homo sapiens CGI-49 protein (LOC51097), mRNA. /FEA=mRNA /GEN=LOC51097 /PROD=CGI-49 protein /DB_XREF=gi:7705766 /UG=Hs.238126 CGI-49 protein /FL=gb:AF151807.1 gb:NM_016002.1	NM_016002		NP_057086
201833_at	0.034854	gb:NM_001527.1 /DEF=Homo sapiens histone deacetylase 2 (HDAC2), mRNA. /FEA=mRNA /GEN=HDAC2 /PROD=histone deacetylase 2 /DB_XREF=gi:4557640 /UG=Hs.3352 histone deacetylase 2 /FL=gb:U31814.1 gb:NM_001527.1	NM_001527		NP_001518
201840_at	7.25E-04	gb:NM_006156.1 /DEF=Homo sapiens neural precursor cell expressed, developmentally down-regulated 8 (NEDD8), mRNA. /FEA=mRNA /GEN=NEDD8 /PROD=neural precursor cell expressed, developmentallydown-regulated 8 /DB_XREF=gi:5453759 /UG=Hs.75512 neural precursor cell expressed, developmentally down-regulated 8 /FL=gb:D23662.1 gb:NM_006156.1	NM_006156		NP_006147

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201845_s_at	7.63E-04	gb:AB029551.1 /DEF=Homo sapiens YEAF1 mRNA for YY1 and E4TF1 associated factor 1, complete cds. /FEA=mRNA /GEN=YEAF1 /PROD=YY1 and E4TF1 associated factor 1 /DB_XREF=gi:6714542 /UG=Hs.7910 RING1 and YY1 binding protein /FL=gb:AF179286.1 gb:AB029551.1 gb:NM_012234.1	AB029551		NP_036366
201846_s_at	0.013527	gb:NM_012234.1 /DEF=Homo sapiens RING1 and YY1 binding protein (RYBP), mRNA. /FEA=mRNA /GEN=RYBP /PROD=RING1 and YY1 binding protein. /DB_XREF=gi:6912639 /UG=Hs.7910 RING1 and YY1 binding protein /FL=gb:AF179286.1 gb:AB029551.1 gb:NM_012234.1	NM_012234		NP_036366
201850_at	0.027672	gb:NM_001747.1 /DEF=Homo sapiens capping protein (actin filament), gelsolin-like (CAPG), mRNA. /FEA=mRNA /GEN=CAPG /PROD=capping protein (actin filament), gelsolin-like /DB_XREF=gi:4502560 /UG=Hs.82422 capping protein (actin filament), gelsolin-like. /FL=gb:BC000728.1 gb:M94345.1 gb:NM_001747.1	NM_001747		NP_001738

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
201857_at	0.047555	gb:NM_016107.1 /DEF=Homo sapiens M-phase phosphoprotein homolog (LOC51663), mRNA. /FEA=mRNA /GEN=LOC51663 /PROD=M-phase phosphoprotein homolog /DB_XREF=gi:7706372 /UG=Hs.173518 M-phase phosphoprotein homolog /FL=gb:BC000376.1 gb:BC000746.1 gb:AF100742.1 gb:NM_016107.1	NM_016107		NP_057191
201862_s_at	0.003965	gb:NM_004735.1 /DEF=Homo sapiens leucine rich repeat (in FLII) interacting protein 1 (LRRFIP1), mRNA. /FEA=mRNA /GEN=LRRFIP1 /PROD=leucine rich repeat (in FLII) interacting protein 1 /DB_XREF=gi:4758689 /UG=Hs.326159 leucine rich repeat (in FLII) interacting protein 1 /FL=gb:NM_004735.1	NM_004735		NP_004726
201863_at	0.003373	gb:NM_014077.1 /DEF=Homo sapiens DKFZP586O0120 protein (DKFZP586O0120), mRNA. /FEA=mRNA /GEN=DKFZP586O0120 /PROD=DKFZP586O0120 protein /DB_XREF=gi:7661695 /UG=Hs.4766 DKFZP586O0120 protein /FL=gb:BC000639.1 gb:AL050157.1 gb:AF151902.1 gb:NM_014077.1	NM_014077		NP_054796
201865_x_at	0.007591	nuclear receptor subfamily 3, group C, member 1 (glucocorticoid receptor)	AI432196	Hs.75772	NP_000167

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201866_s_at	0.003965	gb:NM_000176.1 /DEF=Homo sapiens nuclear receptor subfamily 3, group C, member 1 (NR3C1), mRNA. /FEA=mRNA /GEN=NR3C1 /PROD=nuclear receptor subfamily 3, group C, member 1 /DB_XREF=gi:4504132 /UG=Hs.75772 nuclear receptor subfamily 3, group C, member 1 /FL=gb:M10901.1 gb:NM_000176.1	NM_000176		NP_000167
201867_s_at	0.03065	Consensus includes gb:AW968555 /FEA=EST /DB_XREF=gi:8158396 /DB_XREF=est:EST380631 /UG=Hs.76536 transducin (beta)-like 1 /FL=gb:NM_005647.1	NM_005647		NP_005638
201879_at	0.049282	Consensus includes gb:AI694332 /FEA=EST /DB_XREF=gi:4971672 /DB_XREF=est:wd45e11.x1 /CLONE=IMAGE:2331116 /UG=Hs.181461 ariadne (Drosophila) homolog, ubiquitin-conjugating enzyme E2-binding protein, 1 /FL=gb:AF072832.1 gb:NM_005744.2	NM_005744		NP_005735
201881_s_at	0.014603	gb:NM_005744.2 /DEF=Homo sapiens ariadne (Drosophila) homolog, ubiquitin-conjugating enzyme E2-binding protein, 1 (ARIH1), mRNA. /FEA=mRNA /GEN=ARIH1 /PROD=ariadne homolog /DB_XREF=gi:9966762 /UG=Hs.181461 ariadne (Drosophila) homolog, ubiquitin-conjugating enzyme E2-binding protein, 1 /FL=gb:AF072832.1 gb:NM_005744.2	NM_005744		NP_005735

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
201886_at	0.025377	gb:NM_025230.1 /DEF=Homo sapiens hypothetical protein PRO2389 (PRO2389), mRNA. /FEA=mRNA /GEN=PRO2389 /PROD=hypothetical protein PRO2389 /DB_XREF=gi:13489107 /UG=Hs.283976 hypothetical protein PRO2389 /FL=gb:AF130070.1 gb:AF267858.1 gb:NM_025230.1	NM_025230		NP_079506
201890_at	0.043634	Consensus includes gb:BE966236 /FEA=EST /DB_XREF=gi:11771437 /DB_XREF=est:601660172R1 /CLONE=IMAGE:3905920 /UG=Hs.75319 ribonucleotide reductase M2 polypeptide /FL=gb:NM_001034.1	NM_001034		NP_001025
201892_s_at	0.009189	gb:NM_000884.1 /DEF=Homo sapiens IMP (inosine monophosphate) dehydrogenase 2 (IMPDH2), mRNA. /FEA=mRNA /GEN=IMPDH2 /PROD=IMP (inosine monophosphate) dehydrogenase 2 /DB_XREF=gi:4504688 /UG=Hs.75432 IMP (inosine monophosphate) dehydrogenase 2 /FL=gb:J04208.1 gb:NM_000884.1	NM_000884		NP_000875
201894_s_at	7.76E-04	gb:NM_001920.1 /DEF=Homo sapiens decorin (DCN), mRNA. /FEA=mRNA /GEN=DCN /PROD=decorin /DB_XREF=gi:4503270 /UG=Hs.76152 decorin /FL=gb:M14219.1 gb:NM_001920.1 gb:AF138300.1 gb:AF138304.1	NM_001920		NP_598014

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201895_at	0.012147	gb:NM_001654.1 /DEF=Homo sapiens v-raf murine sarcoma 3611 viral oncogene homolog 1 (ARAF1), mRNA. /FEA=mRNA /GEN=ARAF1 /PROD=v-raf murine sarcoma 3611 viral oncogene homolog1 /DB_XREF=gi:4502192 /UG=Hs.77183 v-raf murine sarcoma 3611 viral oncogene homolog 1 /FL=gb:BC002466.1 gb:NM_001654.1	NM_001654		NP_001645
201897_s_at	0.004515	gb:NM_001826.1 /DEF=Homo sapiens CDC28 protein kinase 1 (CKS1), mRNA. /FEA=mRNA /GEN=CKS1 /PROD=CDC28 protein kinase 1 /DB_XREF=gi:4502856 /UG=Hs.77550 CDC28 protein kinase 1 /FL=gb:BC001425.1 gb:AF274941.1 gb:AF279897.1 gb:NM_001826.1	NM_001826		NP_001817
201898_s_at	7.25E-04	ubiquitin-conjugating enzyme E2A (RAD6 homolog)	AI126625	Hs.80612	NP_003327
201899_s_at	0.007174	gb:NM_003336.1 /DEF=Homo sapiens ubiquitin-conjugating enzyme E2A (RAD6 homolog) (UBE2A), mRNA. /FEA=mRNA /GEN=UBE2A /PROD=ubiquitin-conjugating enzyme E2A (RAD6 homolog) /DB_XREF=gi:4507768 /UG=Hs.80612 ubiquitin-conjugating enzyme E2A (RAD6 homolog) /FL=gb:M74524.1 gb:NM_003336.1	NM_003336		NP_003327

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCanc r					
Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201901_s_at	0.013527	Consensus includes gb:Z14077.1 /DEF=H.sapiens mRNA for YY1NF-E1 protein. /FEA=mRNA /PROD=YY1 NF-E1 /DB_XREF=gi:38010 /UG=Hs.97496 YY1 transcription factor /FL=gb:M77698.1 gb:M76541.1 gb:NM_003403.2	NM_003403		NP_003394
201903_at	0.001763	gb:NM_003365.1 /DEF=Homo sapiens ubiquinol-cytochrome c reductase core protein I (UQCRC1), mRNA. /FEA=mRNA /GEN=UQCRC1 /PROD=ubiquinol-cytochrome c reductase core protein I /DB_XREF=gi:4507840 /UG=Hs.119251 ubiquinol-cytochrome c reductase core protein I /FL=gb:L16842.1 gb:NM_003365.1 gb:D26485.1	NM_003365		NP_003356
201912_s_at	7.63E-04	gb:NM_002094.1 /DEF=Homo sapiens G1 to S phase transition 1 (GSPT1), mRNA. /FEA=mRNA /GEN=GSPT1 /PROD=G1 to S phase transition 1 /DB_XREF=gi:4504166 /UG=Hs.2707 G1 to S phase transition 1 /FL=gb:NM_002094.1	NM_002094		NP_002085
201913_s_at	0.048412	gb:NM_025233.1 /DEF=Homo sapiens nucleotide binding protein (NBP), mRNA. /FEA=mRNA /GEN=NBP /PROD=nucleotide binding protein /DB_XREF=gi:13376837 /UG=Hs.296422 nucleotide binding protein /FL=gb:NM_025233.1 gb:AF208536.1	NM_025233		NP_079509

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201914_s_at	0.010975	Consensus includes gb:AK001465.1 /DEF=Homo sapiens cDNA FLJ10603 fis, clone NT2RP2005012, highly similar to Homo sapiens mRNA for SEC63 protein. /FEA=mRNA /DB_XREF=gi:7022740 /UG=Hs.31575 SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like /FL=gb:AF100141.1 gb:NM_007214.1	AK001465		NP_009145
201915_at	0.018591	Consensus includes gb:AI806665 /FEA=EST /DB_XREF=gi:5393231 /DB_XREF=est:wf35c05.x1 /CLONE=IMAGE:2357576 /UG=Hs.31575 SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like /FL=gb:AF100141.1 gb:NM_007214.1	AK001465		NP_009145
201916_s_at	0.002109	gb:NM_007214.1 /DEF=Homo sapiens SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like (SEC63L), mRNA. /FEA=mRNA /GEN=SEC63L /PROD=SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like /DB_XREF=gi:6005871 /UG=Hs.31575 SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like /FL=gb:AF100141.1 gb:NM_007214.1	NM_007214		NP_009145
201917_s_at	6.66E-04	hypothetical protein FLJ10618	AI694452	Hs.408258	NP_060625
201918_at	0.042466	hypothetical protein FLJ10618	AI694452	Hs.408258	NP_060625

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
201920_at	0.034296	gb:NM_005415.2 /DEF=Homo sapiens solute carrier family 20 (phosphate transporter), member 1 (SLC20A1), mRNA. /FEA=mRNA /GEN=SLC20A1 /PROD=solute carrier family 20 (phosphatetransporter), member 1 /DB_XREF=gi:7382462 /UG=Hs.78452 solute carrier family 20 (phosphate transporter), member 1 /FL=gb:L20859.1 gb:NM_005415.2	NM_005415		NP_005406
201921_at	0.033358	gb:NM_004125.1 /DEF=Homo sapiens guanine nucleotide binding protein 10 (GNG10), mRNA. /FEA=mRNA /GEN=GNG10 /PROD=guanine nucleotide binding protein 10 /DB_XREF=gi:4758445 /UG=Hs.79126 guanine nucleotide binding protein 10 /FL=gb:NM_004125.1 gb:U31383.1	NM_004125		NP_004116
201925_s_at	6.66E-04	gb:NM_000574.1 /DEF=Homo sapiens decay accelerating factor for complement (CD55, Cromer blood group system) (DAF), mRNA. /FEA=mRNA /GEN=DAF /PROD=decay accelerating factor for complement (CD55, Cromer blood group system) /DB_XREF=gi:10835142 /UG=Hs.1369 decay accelerating factor for complement (CD55, Cromer blood group system) /FL=gb:NM_000574.1 gb:BC001288.1 gb:M31516.1	NM_000574		NP_000565

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
201926_s_at	7.25E-04	gb:BC001288.1 /DEF=Homo sapiens, Similar to decay accelerating factor for complement (CD55, Cromer blood group system), clone MGC:5192, mRNA, complete cds. /FEA=mRNA /PROD=Similar to decay accelerating factor for complement (CD55, Cromer blood group system) /DB_XREF=gi:12654888 /UG=Hs.1369 decay accelerating factor for complement (CD55, Cromer blood group system) /FL=gb:NM_000574.1 gb:BC001288.1 gb:M31516.1	BC001288		NP_000565
201928_at	0.026836	Consensus includes gb:AA194254 /FEA=EST /DB_XREF=gi:1783969 /DB_XREF=est:zr39d09:s1 /CLONE=IMAGE:665777 /UG=Hs.152151 plakophilin 4 /FL=gb:NM_003628.2	NM_003628		NP_003619
201929_s_at	0.007984	gb:NM_003628.2 /DEF=Homo sapiens plakophilin 4 (PKP4), mRNA: /FEA=mRNA /GEN=PKP4 /PROD=plakophilin 4 /DB_XREF=gi:6806894 /UG=Hs.152151 plakophilin 4 /FL=gb:NM_003628.2	NM_003628		NP_003619
201930_at	0.001916	gb:NM_005915.2 /DEF=Homo sapiens minichromosome maintenance deficient (mis5, S. pombe) 6 (MCM6), mRNA: /FEA=mRNA /GEN=MCM6 /PROD=minichromosome maintenance deficient (mis5, S.pombe) 6 /DB_XREF=gi:7427518 /UG=Hs.155462 minichromosome maintenance deficient (mis5, S. pombe) 6 /FL=gb:U46838.1 gb:D84557.1 gb:NM_005915.2	NM_005915		NP_005906

Genes Corresponding To Differ ntially Expressed Genes in Figure 18 - BladderCancer					
Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201936_s_at	0.005083	gb:NM_003760.2 /DEF=Homo sapiens eukaryotic translation initiation factor 4 gamma, 3 (EIF4G3), mRNA. /FEA=mRNA /GEN=EIF4G3 /PROD=eukaryotic translation initiation factor 4gamma, 3 /DB_XREF=gi:10092600 /UG=Hs.25732 eukaryotic translation initiation factor 4 gamma, 3 /FL=gb:NM_003760.2 gb:AF012072.2	NM_003760		NP_003751
201940_at	0.002621	carboxypeptidase D	AA897514	Hs.5057	NP_001295
201941_at	0.002621	carboxypeptidase D	AA897514	Hs.5057	NP_001295
201944_at	7.76E-04	gb:NM_000521.2 /DEF=Homo sapiens hexosaminidase B (beta polypeptide) (HEXB), mRNA. /FEA=mRNA /GEN=HEXB /PROD=hexosaminidase B preproprotein /DB_XREF=gi:13128866 /UG=Hs.51043 hexosaminidase B (beta polypeptide) /FL=gb:NM_000521.2 gb:M19735.1	NM_000521		NP_000512
201949_x_at	0.030968	capping protein (actin filament) muscle Z-line, beta	AL572341	Hs.333417	NP_004921
201951_at	0.027066	Consensus includes gb:BF242905 /FEA=EST /DB_XREF=gi:11156833 /DB_XREF=est:601877949F1 /CLONE=IMAGE:4106028 /UG=Hs.10247 activated leucocyte cell adhesion molecule /FL=gb:NM_001627.1 gb:L38608.1	NM_001627		NP_001618

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
201954_at	0.005574	gb:NM_005720.1 /DEF=Homo sapiens actin related protein 23 complex, subunit 1A (41 kD) (ARPC1B), mRNA. /FEA=mRNA /GEN=ARPC1B /PROD=actin related protein 23 complex, subunit 1A(41 kD) /DB_XREF=gi:5031600 /UG=Hs.11538 actin related protein 23 complex, subunit 1A (41 kD) /FL=gb:BC002562.1 gb:AF006084.1 gb:NM_005720.1	NM_005720		NP_005711
201959_s_at	0.006756	KIAA0916 protein	AA488899	Hs.151411	NP_055872
201963_at	0.010197	gb:NM_021122.2 /DEF=Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 2 (FACL2), mRNA. /FEA=mRNA /GEN=FACL2 /PROD=long-chain fatty-acid-Coenzyme A ligase 2. /DB_XREF=gi:12669906 /UG=Hs.154890 fatty-acid-Coenzyme A ligase, long-chain 2 /FL=gb:NM_021122.2 gb:D10040.1	NM_021122		NP_066945
201968_s_at	7.76E-04	gb:NM_002633.1 /DEF=Homo sapiens phosphoglucomutase 1 (PGM1), mRNA. /FEA=mRNA /GEN=PGM1 /PROD=phosphoglucomutase 1 /DB_XREF=gi:4505764 /UG=Hs.1869 phosphoglucomutase 1 /FL=gb:BC001756.1 gb:M83088.1 gb:NM_002633.1	NM_002633		NP_002624

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201972_at	0.018248	gb:AF113129.1 /DEF=Homo sapiens vacuolar ATPase isoform VA68 mRNA, complete cds. /FEA=mRNA /PROD=vacuolar ATPase isoform VA68 /DB_XREF=gi:6523820 /UG=Hs.281866 ATPase, H+ transporting, lysosomal (vacuolar proton pump); alpha polypeptide, 70kD, isoform 1 /FL=gb:L09235.1 gb:NM_001690.1 gb:AF113129.1	AF113129		NP_001681
201973_s_at	0.014372	CGI-43 protein	AL550875	Hs.289112	NP_056437
201975_at	0.002446	gb:NM_002956.1 /DEF=Homo sapiens restin (Reed-Steinberg cell-expressed intermediate filament-associated protein) (RSN), mRNA. /FEA=mRNA /GEN=RSN /PROD=restin (Reed-Steinberg cell-expressed intermediate filament-associated protein) /DB_XREF=gi:4506750 /UG=Hs.31638 restin (Reed-Steinberg cell-expressed intermediate filament-associated protein) /FL=gb:NM_002956.1	NM_002956		NP_002947
201980_s_at	0.010713	gb:NM_012425.2 /DEF=Homo sapiens Ras suppressor protein 1 (RSU1), mRNA. /FEA=mRNA /GEN=RSU1 /PROD=ras suppressor protein 1 /DB_XREF=gi:10800408 /UG=Hs.75551 Homo sapiens Ras suppressor protein 1 (RSU1), mRNA /FL=gb:NM_012425.2	NM_012425		NP_036557

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201985_at	7.63E-04	gb:NM_014846.1 /DEF=Homo sapiens KIAA0196 gene product (KIAA0196), mRNA. /FEA=mRNA /GEN=KIAA0196 /PROD=KIAA0196 gene product /DB_XREF=gi:7661987 /UG=Hs.8294 KIAA0196 gene product /FL=gb:D83780.1 gb:NM_014846.1	NM_014846		NP_055661
201995_at	7.25E-04	gb:NM_000127.1 /DEF=Homo sapiens exostoses (multiple) 1 (EXT1), mRNA. /FEA=mRNA /GEN=EXT1 /PROD=exostoses (multiple) 1 /DB_XREF=gi:4557570 /UG=Hs.184161 exostoses (multiple) 1 /FL=gb:BC001174.1 gb:NM_000127.1	NM_000127		NP_000118
201997_s_at	0.029624	gb:NM_015001.1 /DEF=Homo sapiens KIAA0929 protein Msx2 interacting nuclear target (MINT) homolog (KIAA0929), mRNA. /FEA=mRNA /GEN=KIAA0929 /PROD=KIAA0929 protein Msx2 interacting nuclear target(MINT) homolog /DB_XREF=gi:7657266 /UG=Hs.184245 KIAA0929 protein Msx2 interacting nuclear target (MINT) homolog /FL=gb:NM_015001.1	NM_015001		NP_055816

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201999_s_at	0.029279	gb:NM_006519.1 /DEF=Homo sapiens t-complex-associated-testis-expressed 1-like 1 (TCTEL1), mRNA. /FEA=mRNA /GEN=TCTEL1 /PROD=t-complex-associated-testis-expressed 1-like 1 /DB_XREF=gi:5730084 /UG=Hs.266940 t-complex-associated-testis-expressed 1-like 1 /FL=gb:U56255.1 gb:D50663.1 gb:NM_006519.1	NM_006519		NP_006510
202000_at	0.031551	gb:BC002772.1 /DEF=Homo sapiens, NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14), clone MGC:3686; mRNA, complete cds. /FEA=mRNA /PROD=NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14) /DB_XREF=gi:12803858 /UG=Hs.274416 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14) /FL=gb:BC002772.1 gb:AF047182.1 gb:NM_002490.1	BC002772		NP_002481
202006_at	7.63E-04	gb:NM_002835.1 /DEF=Homo sapiens protein tyrosine phosphatase, non-receptor type 12 (PTPN12), mRNA. /FEA=mRNA /GEN=PTPN12 /PROD=protein tyrosine phosphatase, non-receptor type 12 /DB_XREF=gi:4506286 /UG=Hs.62 protein tyrosine phosphatase, non-receptor type 12 /FL=gb:D13380.1 gb:M93425.1 gb:NM_002835.1	NM_002835		NP_002826

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
202009_at	0.049173	gb:NM_007284.1 /DEF=Homo sapiens protein tyrosine kinase 9-like (A6-related protein) (PTK9L), mRNA. /FEA=mRNA /GEN=PTK9L /PROD=protein tyrosine kinase 9-like (A6-related protein) /DB_XREF=gi:6005845 /UG=Hs.6780 protein tyrosine kinase 9-like (A6-related protein) /FL=gb:BC000327.1 gb:BC003161.1 gb:NM_007284.1 gb:AL136773.1	NM_007284		NP_009215
202010_s_at	0.001852	gb:NM_021188.1 /DEF=Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA. /FEA=mRNA /GEN=LOC57862 /PROD=clones 23667 and 23775 zinc finger protein /DB_XREF=gi:10863994 /UG=Hs.7137 clones 23667 and 23775 zinc finger protein /FL=gb:NM_021188.1 gb:U90919.1	NM_021188		NP_067011
202018_s_at	0.006178	gb:NM_002343.1 /DEF=Homo sapiens lactotransferrin (LTF), mRNA. /FEA=mRNA /GEN=LTF /PROD=lactotransferrin /DB_XREF=gi:4505042 /UG=Hs.105938 lactotransferrin /FL=gb:AF332168.1 gb:M93150.1 gb:M83202.1 gb:NM_002343.1	NM_002343		NP_002334

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unig ne Accession No.	Protein Accession No.
202025_x_at	0.007652	gb:NM_001607.2 /DEF=Homo sapiens acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase) (ACAA1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=ACAA1 /PROD=acetyl-Coenzyme A acyltransferase 1 /DB_XREF=gi:6598316 /UG=Hs.166160 acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase) /FL=gb:BC000635.1 gb:NM_001607.2	NM_001607		NP_001598
202026_at	0.001213	gb:NM_003002.1 /DEF=Homo sapiens succinate dehydrogenase complex, subunit D, integral membrane protein (SDHD), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=SDHD /PROD=succinate dehydrogenase complex, subunit Dprecursor /DB_XREF=gi:4506864 /UG=Hs.168289 succinate dehydrogenase complex, subunit D, integral membrane protein /FL=gb:BC005263.1 gb:AB006202.1 gb:NM_003002.1	NM_003002		NP_002993
202027_at	0.032723	gb:NM_012264.1 /DEF=Homo sapiens chromosome 22 open reading frame 5 (C22ORF5), mRNA. /FEA=mRNA /GEN=C22ORF5 /PROD=chromosome 22 open reading frame 5 /DB_XREF=gi:7110634 /UG=Hs.182626 chromosome 22 open reading frame 5 /FL=gb:NM_012264.1	NM_012264		NP_036396

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202028_s_at	0.03947	Consensus includes gb:BC000603.1 /DEF=Homo sapiens, ribosomal protein L38, clone MGC:1637, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein L38 /DB_XREF=gi:12653644 /UG=Hs.2017 ribosomal protein L38 /FL=gb:BC000603.1 gb:NM_000999.1	BC000603		NP_000990
202032_s_at	0.012835	gb:NM_006122.1 /DEF=Homo sapiens mannosidase, alpha, class 2A, member 2 (MAN2A2), mRNA. /FEA=mRNA /GEN=MAN2A2 /PROD=mannosidase, alpha, class 2A, member 2 /DB_XREF=gi:5540099 /UG=Hs.295605 mannosidase, alpha, class 2A, member 2 /FL=gb:NM_006122.1 gb:L28821.1	NM_006122		NP_006113
202040_s_at	7.76E-04	gb:NM_005056.1 /DEF=Homo sapiens retinoblastoma-binding protein 2 (RBBP2), mRNA. /FEA=mRNA /GEN=RBBP2 /PROD=retinoblastoma-binding protein 2 /DB_XREF=gi:4826967 /UG=Hs.76272 retinoblastoma-binding protein 2 /FL=gb:NM_005056.1	NM_005056		NP_005047

Genes Corresponding To Differ ntially Expr ss d G nes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Acc ssion No.
202041_s_at	6.66E-04	gb:NM_004214.3 /DEF=Homo sapiens fibroblast growth factor (acidic) intracellular binding protein (FIBP), mRNA. /FEA=mRNA /GEN=FIBP /PROD=fibroblast growth factor (acidic) intracellular binding protein /DB_XREF=gi:7262377 /UG=Hs.7768 fibroblast growth factor (acidic) intracellular binding protein /FL=gb:AF171944.1 gb:AF171945.1 gb:AF171946.1 gb:AF010187.2 gb:NM_004214.3	NM_004214		NP_004205
202047_s_at	0.01063	chromobox homolog 6	AI458128	Hs.107374	NP_055107
202055_at	0.003489	Consensus includes gb:AA652173 /FEA=EST /DB_XREF=gi:2583825 /DB_XREF=est:ns51f05.s1 /CLONE=IMAGE:1187169 /UG=Hs.169149 karyopherin alpha 1 (importin alpha 5) /FL=gb:BC002374.1 gb:BC003009.1 gb:NM_002264.1	NM_002264		NP_002255
202059_s_at	7.25E-04	gb:NM_002264.1 /DEF=Homo sapiens karyopherin alpha 1 (importin alpha 5) (KPNA1), mRNA. /FEA=mRNA /GEN=KPNA1 /PROD=karyopherin alpha 1 /DB_XREF=gi:4504894 /UG=Hs.169149 karyopherin alpha 1 (importin alpha 5) /FL=gb:BC002374.1 gb:BC003009.1 gb:NM_002264.1	NM_002264		NP_002255
202061_s_at	7.63E-04	sel-1 suppressor of lin-12-like (C. elegans)	AI927770	Hs.181300	NP_005056

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
202066_at	0.027176	Consensus includes gb:AA195259 /FEA=EST /DB_XREF=gi:1784959 /DB_XREF=est:zr36g01:s1 /CLONE=IMAGE:665520 /UG=Hs.183648 protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 1 /FL=gb:NM_003626.1 gb:U22816.1	NM_003626		NP_803172
202068_s_at	0.018398	gb:NM_000527.2 /DEF=Homo sapiens low density lipoprotein receptor (familial hypercholesterolemia) (LDLR), mRNA. /FEA=mRNA /GEN=LDLR /PROD=low density lipoprotein receptor precursor /DB_XREF=gi:8051613 /UG=Hs.213289 low density lipoprotein receptor (familial hypercholesterolemia) /FL=gb:NM_000527.2	NM_000527		NP_000518
202079_s_at	0.017638	KIAA1042 protein	AI633774	Hs.6705	NP_055780
202080_s_at	8.74E-04	gb:NM_014965.1 /DEF=Homo sapiens KIAA1042 protein (KIAA1042), mRNA. /FEA=mRNA /GEN=KIAA1042 /PROD=KIAA1042 protein /DB_XREF=gi:7662457 /UG=Hs.6705 KIAA1042 protein /FL=gb:AB028965.1 gb:NM_014965.1	NM_014965		NP_055780
202082_s_at	0.014875	Consensus includes gb:AV748469 /FEA=EST /DB_XREF=gi:10906317 /DB_XREF=est:AV748469 /CLONE=NPCAXA07 /UG=Hs.75232 SEC14 (S. cerevisiae)-like 1 /FL=gb:D67029.1 gb:NM_003003.1	NM_003003		NP_002994

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202087_s_at	0.011409	gb:NM_001912.1 /DEF=Homo sapiens cathepsin L (CTSL), mRNA. /FEA=mRNA /GEN=CTSL /PROD=cathepsin L /DB_XREF=gi:4503154 /UG=Hs.78056 cathepsin L /FL=gb:NM_001912.1	NM_001912		NP_666023
202088_at	0.025231	LIV-1 protein, estrogen regulated	AI635449	Hs.79136	NP_036451
202089_s_at	0.005347	gb:NM_012319.2 /DEF=Homo sapiens LIV-1 protein, estrogen regulated (LIV-1), mRNA. /FEA=mRNA /GEN=LIV-1 /PROD=LIV-1 protein, estrogen regulated /DB_XREF=gi:12751474 /UG=Hs.79136 LIV-1 protein, estrogen regulated /FL=gb:U41060.2 gb:NM_012319.2	NM_012319		NP_036451
202090_s_at	0.002749	gb:NM_006830.1 /DEF=Homo sapiens ubiquinol-cytochrome c reductase (6.4kD) subunit (UQCR), mRNA. /FEA=mRNA /GEN=UQCR /PROD=ubiquinol-cytochrome c reductase (6.4kD) subunit /DB_XREF=gi:5803216 /UG=Hs.8372 ubiquinol-cytochrome c reductase (6.4kD) subunit /FL=gb:D55636.1 gb:NM_006830.1	NM_006830		NP_006821
202093_s_at	0.048412	gb:NM_019088.1 /DEF=Homo sapiens hypothetical protein F23149_1 (F23149_1), mRNA. /FEA=mRNA /GEN=F23149_1 /PROD=hypothetical protein F23149_1 /DB_XREF=gi:9506582 /UG=Hs.152894 hypothetical protein F23149_1 /FL=gb:BC000017.1 gb:NM_019088.1	NM_019088		NP_061961

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202097_at	0.005652	gb:NM_005124.1 /DEF=Homo sapiens nucleoporin 153kD (NUP153), mRNA. /FEA=mRNA /GEN=NUP153 /PROD=nucleoporin 153kD /DB_XREF=gi:4826871 /UG=Hs.211608 nucleoporin 153kD /FL=gb:NM_005124.1	NM_005124		NP_005115
202098_s_at	0.022439	gb:NM_001535.1 /DEF=Homo sapiens HMT1 (hnRNP methyltransferase, S. cerevisiae)-like 1 (HRMT1L1), mRNA. /FEA=mRNA /GEN=HRMT1L1 /PROD=HMT1 (hnRNP methyltransferase, S. cerevisiae)-like 1 /DB_XREF=gi:4504494 /UG=Hs.235887 HMT1 (hnRNP methyltransferase, S. cerevisiae)-like 1 /FL=gb:U80213.1 gb:NM_001535.1	NM_001535		NP_001526
202100_at	0.001406	v-rat simian leukemia viral oncogene homolog B (ras related; GTP binding protein)	BG169673	Hs.348024	NP_002872
202102_s_at	0.012147	bromodomain containing 4	BF718610	Hs.278675	NP_490597
202105_at	0.001641	gb:NM_001551.1 /DEF=Homo sapiens immunoglobulin (CD79A) binding protein 1 (IGBP1), mRNA. /FEA=mRNA /GEN=IGBP1 /PROD=immunoglobulin-binding protein 1 /DB_XREF=gi:4557662 /UG=Hs.3631 immunoglobulin (CD79A) binding protein 1 /FL=gb:BC004137.1 gb:NM_001551.1	NM_001551		NP_001542

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202113_s_at	0.049392	gb:AF043453.1 /DEF=Homo sapiens sorting nexin 2 (SNX2) mRNA, complete cds. /FEA=mRNA /GEN=SNX2 /PROD=sorting nexin 2 /DB_XREF=gi:2827433 /UG=Hs.11183 sorting nexin 2 /FL=gb:BC003382.1 gb:AF043453.1 gb:AF065482.1 gb:NM_003100.1	AF043453		
202119_s_at	6.66E-04	gb:NM_003909.1 /DEF=Homo sapiens copine III (CPNE3), mRNA. /FEA=mRNA /GEN=CPNE3 /PROD=copine III /DB_XREF=gi:4503014 /UG=Hs.14158 copine III /FL=gb:AB014536.1 gb:NM_003909.1 gb:AF077226.2	NM_003909		NP_003900
202120_x_at	0.040104	gb:NM_004069.2 /DEF=Homo sapiens adaptor-related protein complex 2, sigma 1 subunit (AP2S1), transcript variant AP17, mRNA. /FEA=mRNA /GEN=AP2S1 /PROD=adaptor-related protein complex 2, sigma 1 subunit, isoform AP17 /DB_XREF=gi:11038644 /UG=Hs.119591 adaptor-related protein complex 2, sigma 1 subunit /FL=gb:NM_004069.2	NM_004069		NP_067586
202121_s_at	0.00121	gb:NM_014453.1 /DEF=Homo sapiens putative breast adenocarcinoma marker (32kD) (BC-2), mRNA. /FEA=mRNA /GEN=BC-2 /PROD=putative breast adenocarcinoma marker (32kD) /DB_XREF=gi:7656921 /UG=Hs.12107 putative breast adenocarcinoma marker (32kD) /FL=gb:AF042384.1 gb:NM_014453.1	NM_014453		NP_055268

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202122_s_at	0.041179	gb:NM_005817.1 /DEF=Homo sapiens cargo selection protein (mannose 6 phosphate receptor binding protein) (TIP47), mRNA. /FEA=mRNA /GEN=TIP47 /PROD=cargo selection protein (mannose 6 phosphatereceptor binding protein) /DB_XREF=gi:5032182 /UG=Hs.140452 cargo selection protein (mannose 6 phosphate receptor binding protein) /FL=gb:AF057140.1 gb:AF051314.1 gb:AF051315.1 gb:AF055574.1 gb:NM_005817.1	NM_005817		NP_005808
202124_s_at	0.03065	amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3	AV705253	Hs.154248	NP_055864
202125_s_at	0.014263	gb:NM_015049.1 /DEF=Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 (ALS2CR3), mRNA. /FEA=mRNA /GEN=ALS2CR3 /PROD=amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 /DB_XREF=gi:13027379 /UG=Hs.154248 amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 /FL=gb:AB038951.1 gb:NM_015049.1	NM_015049		NP_055864

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202128_at	0.002749	gb:NM_014821.1 /DEF=Homo sapiens KIAA0317 gene product (KIAA0317), mRNA. /FEA=mRNA /GEN=KIAA0317 /PROD=KIAA0317 gene product /DB_XREF=gi:7662051 /UG=Hs.20126 KIAA0317 gene product /FL=gb:AB002315.1 gb:NM_014821.1	NM_014821		NP_055636
202129_s_at	0.006449	sudD suppressor of bimD6 homolog (A. nidulans)	AW006290	Hs.209061	NP_665913
202130_at	7.63E-04	sudD suppressor of bimD6 homolog (A. nidulans)	AW006290	Hs.209061	NP_665913
202131_s_at	7.76E-04	gb:NM_003831.1 /DEF=Homo sapiens sudD (suppressor of bimD6, Aspergillus nidulans) homolog (SUDD); mRNA. /FEA=mRNA /GEN=SUDD /PROD=sudD (suppressor of bimD6, Aspergillus nidulans)homolog /DB_XREF=gi:4507298 /UG=Hs.209061 sudD (suppressor of bimD6, Aspergillus nidulans) homolog /FL=gb:AF013591.1 gb:NM_003831.1	NM_003831		NP_665913
202135_s_at	0.015807	gb:NM_005735.2 /DEF=Homo sapiens ARP1 (actin-related protein 1, yeast) homolog B (centractin beta) (ACTR1B), mRNA. /FEA=mRNA /GEN=ACTR1B /PROD=actin-related protein 1B /DB_XREF=gi:13325060 /UG=Hs.2477 ARP1 (actin-related protein 1, yeast) homolog B (centractin beta) /FL=gb:NM_005735.2 gb:BC004374.1	NM_005735		NP_005726
202136_at	0.024403	adenovirus 5 E1A binding protein	BE250417	Hs.301449	NP_006615

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
202139_at	0.021022	gb:NM_003689.1 /DEF=Homo sapiens aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase) (AKR7A2), mRNA. /FEA=mRNA /GEN=AKR7A2 /PROD=aldo-keto reductase family 7, member A2(aflatoxin aldehyde reductase) /DB_XREF=gi:4502020 /UG=Hs.6980 aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase) /FL=gb:BC004111.1 gb:AF026947.1 gb:NM_003689.1	NM_003689		NP_003680
202140_s_at	0.009329	gb:NM_003992.1 /DEF=Homo sapiens CDC-like kinase 3 (CLK3), transcript variant phclk3, mRNA. /FEA=mRNA /GEN=CLK3 /PROD=CDC-like kinase 3 isoform hclk3 /DB_XREF=gi:4502884 /UG=Hs.73987 CDC-like kinase 3 /FL=gb:BC002555.1 gb:NM_003992.1 gb:L29217.1	NM_003992		NP_003983
202143_s_at	0.020576	gb:NM_006710.1 /DEF=Homo sapiens COP9 homolog (COP9), mRNA. /FEA=mRNA /GEN=COP9 /PROD=COP9 homolog /DB_XREF=gi:5729778 /UG=Hs.75193 COP9 homolog /FL=gb:BC003090.1 gb:U51205.1 gb:NM_006710.1	NM_006710		NP_006701
202155_s_at	7.63E-04	gb:NM_005085.1 /DEF=Homo sapiens nucleoporin 214kD (CAIN) (NUP214), mRNA. /FEA=mRNA /GEN=NUP214 /PROD=nucleoporin 214kD (CAIN) /DB_XREF=gi:4826873 /UG=Hs.170285 nucleoporin 214kD (CAIN) /FL=gb:NM_005085.1	NM_005085		NP_705906

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202160_at	0.003875	gb:NM_004380.1 /DEF=Homo sapiens CREB binding protein (Rubinstein-Taybi syndrome) (CREBBP), mRNA. /FEA=mRNA /GEN=CREBBP /PROD=CREB binding protein /DB_XREF=gi:4758055 /UG=Hs.23598 CREB binding protein (Rubinstein-Taybi syndrome) /FL=gb:U47741.1 gb:NM_004380.1	NM_004380		NP_004371
202162_s_at	0.001852	TNFAIP3 interacting protein 1	AI769416	Hs.109281	NP_004770
202163_s_at	0.007591	gb:NM_004779.1 /DEF=Homo sapiens CCR4-NOT transcription complex, subunit 8 (CNOT8), mRNA. /FEA=mRNA /GEN=CNOT8 /PROD=CCR4-NOT transcription complex, subunit 8 /DB_XREF=gi:4758945 /UG=Hs.26703 CCR4-NOT transcription complex, subunit 8 /FL=gb:AF053318.1 gb:NM_004779.1 gb:AL122045.1 gb:AF180476.1	NM_004779		NP_004770
202164_s_at	7.76E-04	gb:AF180476.1 /DEF=Homo sapiens CALIFp (CALIF) mRNA, complete cds. /FEA=mRNA /GEN=CALIF /PROD=CALIFp /DB_XREF=gi:6856208 /UG=Hs.26703 CCR4-NOT transcription complex, subunit 8 /FL=gb:AF053318.1 gb:NM_004779.1 gb:AL122045.1 gb:AF180476.1	AF180476		NP_004770

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202172_at	0.030968	Consensus includes gb:BG035116 /FEA=EST /DB_XREF=gi:12428927 /DB_XREF=est:602324811F1 /CLONE=IMAGE:4412907 /UG=Hs.6557 zinc finger protein 161 /FL=gb:D28118.1 gb:NM_007146.1	NM_007146		NP_009077
202181_at	0.005347	gb:NM_014734.1 /DEF=Homo sapiens KIAA0247 gene product (KIAA0247), mRNA. /FEA=mRNA /GEN=KIAA0247 /PROD=KIAA0247 gene product /DB_XREF=gi:7662019 /UG=Hs.82426 KIAA0247 gene product /FL=gb:D87434.1 gb:NM_014734.1	NM_014734		NP_055549
202182_at	0.003393	gb:NM_021078.1 /DEF=Homo sapiens GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 2 (GCN5L2), mRNA. /FEA=mRNA /GEN=GCN5L2 /PROD=GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 2 /DB_XREF=gi:10835100 /UG=Hs.101067 GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 2 /FL=gb:NM_021078.1	NM_021078		NP_066564

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gen Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
202187_s_at	0.001108	gb:NM_006243.1 /DEF=Homo sapiens protein phosphatase 2, regulatory subunit B (B56), alpha isoform (PPP2R5A), mRNA. /FEA=mRNA /GEN=PPP2R5A /PROD=protein phosphatase 2, regulatory subunit B(B56), alpha isoform /DB_XREF=gi:5453949 /UG=Hs.155079 protein phosphatase 2, regulatory subunit B (B56), alpha isoform /FL=gb:L42373.1 gb:NM_006243.1	NM_006243		NP_006234
202192_s_at	6.66E-04	gb:NM_005890.1 /DEF=Homo sapiens growth arrest-specific 7 (GAS7), transcript variant b, mRNA. /FEA=mRNA /GEN=GAS7 /PROD=growth arrest-specific 7 isoform b /DB_XREF=gi:5360211 /UG=Hs.226133 growth arrest-specific 7 /FL=gb:AB007854.1 gb:NM_005890.1	NM_005890		NP_005881
202193_at	0.010197	gb:NM_005569.2 /DEF=Homo sapiens LIM domain kinase 2 (LIMK2), transcript variant 2a, mRNA. /FEA=mRNA /GEN=LIMK2 /PROD=LIM domain kinase 2 isoform 2a /DB_XREF=gi:8051619 /UG=Hs.278027 LIM domain kinase 2 /FL=gb:D45906.1 gb:NM_005569.2	NM_005569		NP_057952

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202194_at	0.01263	Consensus includes gb:AL117354 /DEF=Human DNA sequence from clone RP5-976O13 on chromosome 1p21.2-22.2 Contains part of the gene for CGI-100 protein, 3 isoforms of the gene for M96 protein, ESTs, STSs, GSSs and a CpG Island /FEA=mRNA_1 /DB_XREF=gi:6822199 /UG=Hs.296155 CGI-100 protein /FL=gb:AF151858.1 gb:NM_016040.1	AL117354		
202197_at	7.76E-04	gb:NM_021090.1 /DEF=Homo sapiens myotubularin related protein 3 (MTMR3), mRNA. /FEA=mRNA /GEN=MTMR3 /PROD=myotubularin related protein 3 /DB_XREF=gi:10835108 /UG=Hs.63302 myotubularin related protein 3 /FL=gb:NM_021090.1 gb:AB002369.1 gb:AF233438.1	NM_021090		NP_694691
202200_s_at	7.63E-04	gb:NM_003137.1 /DEF=Homo sapiens SFRS protein kinase 1 (SRPK1), mRNA. /FEA=mRNA /GEN=SRPK1 /PROD=SFRS protein kinase 1 /DB_XREF=gi:4507218 /UG=Hs.75761 SFRS protein kinase 1 /FL=gb:NM_003137.1 gb:U09564.1	NM_003137		NP_003128
202201_at	7.63E-04	gb:NM_000713.1 /DEF=Homo sapiens biliverdin reductase B (flavin reductase (NADPH)) (BLVRB), mRNA. /FEA=mRNA /GEN=BLVRB /PROD=biliverdin reductase B (flavin reductase(NADPH)) /DB_XREF=gi:4502418 /UG=Hs.76289 biliverdin reductase B (flavin reductase (NADPH)) /FL=gb:D26308.1 gb:NM_000713.1	NM_000713		NP_000704

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202206_at	9.85E-04	Consensus includes gb:AW450363 /FEA=EST /DB_XREF=gi:6991139 /DB_XREF=est:UI-H-BI3-akn-d-02-Q-UI.s1 /CLONE=IMAGE:2734875 /UG=Hs.111554 ADP-ribosylation factor-like 7 /FL=gb:BC001051.1 gb:AB016811.1 gb:NM_005737.2	NM_005737		NP_005728
202207_at	0.007174	Consensus includes gb:BG435404 /FEA=EST /DB_XREF=gi:13341910 /DB_XREF=est:602507678F1 /CLONE=IMAGE:4605066 /UG=Hs.111554 ADP-ribosylation factor-like 7 /FL=gb:BC001051.1 gb:AB016811.1 gb:NM_005737.2	NM_005737		NP_005728
202208_s_at	0.006756	gb:BC001051.1 /DEF=Homo sapiens, ADP-ribosylation factor-like 7, clone MGC:1575, mRNA, complete cds. /FEA=mRNA /PROD=ADP-ribosylation factor-like 7 /DB_XREF=gi:12654450 /UG=Hs.111554 ADP-ribosylation factor-like 7 /FL=gb:BC001051.1 gb:AB016811.1 gb:NM_005737.2	BC001051		NP_005728
202211_at	0.001406	gb:BC005122.1 /DEF=Homo sapiens, ADP-ribosylation factor GTPase activating protein 1, clone MGC:10272, mRNA, complete cds. /FEA=mRNA /PROD=ADP-ribosylation factor GTPase activating protein 1 /DB_XREF=gi:13477296 /UG=Hs.13014 ADP-ribosylation factor GTPase activating protein 1 /FL=gb:BC005122.1 gb:AF111847.1 gb:NM_014570.1	BC005122		NP_055385

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202215_s_at	0.003965	gb:NM_014223.2 /DEF=Homo sapiens nuclear transcription factor Y, gamma (NFYC), mRNA. /FEA=mRNA /GEN=NFYC /PROD=nuclear transcription factor Y, gamma /DB_XREF=gi:11496977 /UG=Hs.168157 nuclear transcription factor Y, gamma /FL=gb:NM_014223.2 gb:D85425.1 gb:BC005003.1 gb:D89986.1	NM_014223		NP_055038
202221_s_at	7.63E-04	AV727101 HTC Homo sapiens cDNA clone HTCBGF05 5', mRNA sequence.	AV727101		
202228_s_at	7.63E-04	gb:NM_017455.1 /DEF=Homo sapiens stromal cell derived factor receptor 1 (SDFR1), transcript variant alpha, mRNA. /FEA=mRNA /GEN=SDFR1 /PROD=stromal cell derived factor receptor 1 isoforma /DB_XREF=gi:9257239 /UG=Hs.6354 stromal cell derived factor receptor 1 /FL=gb:AF109127.1 gb:NM_017455.1	NM_017455		NP_059429
202233_s_at	0.030776	gb:NM_006004.1 /DEF=Homo sapiens ubiquinol-cytochrome c reductase hinge protein (UQCRH), mRNA. /FEA=mRNA /GEN=UQCRH /PROD=ubiquinol-cytochrome c reductase hinge protein /DB_XREF=gi:5174744 /UG=Hs.73818 ubiquinol-cytochrome c reductase hinge protein /FL=gb:BC001426.1 gb:BC001934.1 gb:M36647.1 gb:NM_006004.1	NM_006004		NP_005995

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202239_at	0.003875	gb:NM_006437.2 /DEF=Homo sapiens ADP-ribosyltransferase (NAD ⁺ ; poly (ADP-ribose) polymerase)-like 1 (ADPRTL1), mRNA. /FEA=mRNA /GEN=ADPRTL1 /PROD=poly(ADP-ribosyl)transferase-like 1 /DB_XREF=gi:11496990 /UG=Hs.77225 ADP-ribosyltransferase (NAD ⁺ ; poly (ADP-ribose) polymerase)-like 1 /FL=gb:NM_006437.2 gb:AF057160.1 gb:AF158255.1	NM_006437		NP_006428
202241_at	0.007503	gb:NM_025195.1 /DEF=Homo sapiens phosphoprotein regulated by mitogenic pathways (C8FW), mRNA. /FEA=mRNA /GEN=C8FW /PROD=G-protein-coupled receptor induced protein /DB_XREF=gi:13399327 /UG=Hs.7837 phosphoprotein regulated by mitogenic pathways /FL=gb:AF205437.1 gb:NM_025195.1	NM_025195		NP_079471
202243_s_at	0.001213	gb:NM_002796.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, beta type, 4 (PSMB4), mRNA. /FEA=mRNA /GEN=PSMB4 /PROD=proteasome (prosome, macropain) subunit, betatype, 4 /DB_XREF=gi:4506198 /UG=Hs.89545 proteasome (prosome, macropain) subunit, beta type, 4 /FL=gb:BC000331.1 gb:NM_002796.1 gb:D26600.1	NM_002796		NP_002787

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202244_at	0.024403	gb:NM_002796.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, beta type, 4 (PSMB4), mRNA. /FEA=mRNA /GEN=PSMB4 /PROD=proteasome (prosome, macropain) subunit, betatype, 4 /DB_XREF=gi:4506198 /UG=Hs.89545 proteasome (prosome, macropain) subunit, beta type, 4 /FL=gb:BC000331.1 gb:NM_002796.1 gb:D26600.1	NM_002796		NP_002787
202247_s_at	0.003616	metastasis associated 1	BE561596	Hs.101448	NP_004680
202252_at	7.83E-04	gb:NM_002870.1 /DEF=Homo sapiens RAB13, member RAS oncogene family (RAB13), mRNA. /FEA=mRNA /GEN=RAB13 /PROD=RAB13, member RAS oncogene family /DB_XREF=gi:4506362 /UG=Hs.151536 RAB13, member RAS oncogene family /FL=gb:BC000799.1 gb:NM_002870.1	NM_002870		NP_002861
202257_s_at	0.00534	gb:NM_006110.1 /DEF=Homo sapiens CD2 antigen (cytoplasmic tail)-binding protein 2 (CD2BP2), mRNA. /FEA=mRNA /GEN=CD2BP2 /PROD=CD2 antigen (cytoplasmic tail)-binding protein2 /DB_XREF=gi:5174408 /UG=Hs.202677 CD2 antigen (cytoplasmic tail)-binding protein 2 /FL=gb:BC000495.1 gb:BC001947.1 gb:AF104222.1 gb:NM_006110.1	NM_006110		NP_006101

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
202263_at	7.63E-04	gb:NM_016243.1 /DEF=Homo sapiens cytochrome b5 reductase 1 (B5R.1) (LOC51706), mRNA. /FEA=mRNA /GEN=LOC51706 /PROD=cytochrome b5 reductase 1 (B5R.1) /DB_XREF=gi:7706454 /UG=Hs.289113 cytochrome b5 reductase 1 (B5R.1) /FL=gb:AF169481.1 gb:AF125533.1 gb:NM_016243.1	NM_016243		NP_057327
202269_x_at	0.001835	gb:BC002666.1 /DEF=Homo sapiens, guanylate binding protein 1, interferon-inducible, 67kD, clone MGC:3949, mRNA, complete cds. /FEA=mRNA /PROD=guanylate binding protein 1, interferon-inducible, 67kD /DB_XREF=gi:12803662 /UG=Hs.62661 guanylate binding protein 1; interferon-inducible, 67kD /FL=gb:BC002666.1 gb:M55542.1 gb:NM_002053.1	BC002666		NP_002044
202277_at	0.004519	serine palmitoyltransferase, long chain base subunit 1	AL568804	Hs.90458	
202278_s_at	0.021256	gb:NM_006415.1 /DEF=Homo sapiens serine palmitoyltransferase, long chain base subunit 1 (SPTLC1), mRNA. /FEA=mRNA /GEN=SPTLC1 /PROD=serine palmitoyltransferase, long chain base subunit 1. /DB_XREF=gi:5454083 /UG=Hs.90458 serine palmitoyltransferase, long chain base subunit 1 /FL=gb:NM_006415.1	NM_006415		NP_847894

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
202279_at	0.038202	gb:NM_004894.1 /DEF=Homo sapiens chromosome 14 open reading frame 2 (C14ORF2); mRNA. /FEA=mRNA /GEN=C14ORF2 /PROD=chromosome 14 open reading frame 2 /DB_XREF=gi:4758939 /UG=Hs.109052 chromosome 14 open reading frame 2 /FL=gb:BC000429.1 gb:BC001944.1 gb:AF054175.1 gb:NM_004894.1	NM_004894		NP_004885
202281_at	0.017845	gb:NM_005255.1 /DEF=Homo sapiens cyclin G associated kinase (GAK), mRNA. /FEA=mRNA /GEN=GAK /PROD=cyclin G associated kinase /DB_XREF=gi:4885250 /UG=Hs.153227 cyclin G associated kinase /FL=gb:D88435.1 gb:NM_005255.1	NM_005255		NP_005246
202284_s_at	0.046021	gb:NM_000389.1 /DEF=Homo sapiens cyclin-dependent kinase inhibitor 1A (p21, Cip1) (CDKN1A), mRNA. /FEA=mRNA /GEN=CDKN1A /PROD=cyclin-dependent kinase inhibitor 1A (p21, Cip1) /DB_XREF=gi:11386202 /UG=Hs.179665 cyclin-dependent kinase inhibitor 1A (p21, Cip1) /FL=gb:NM_000389.1 gb:BC000275.1 gb:BC001935.1 gb:U03106.1 gb:L26165.1 gb:L25610.1 gb:U09579.1	NM_000389		NP_510867
202286_s_at	0.001093	GA733-1 protein precursor; Human gastrointestinal tumor-associated antigen GA733-1 protein gene, complete cds, clone 05516.	NM_002353		NP_002344

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
202290_at	0.013678	gb:NM_014891.1 /DEF=Homo sapiens PDGF associated protein (HASPP28), mRNA. /FEA=mRNA /GEN=HASPP28 /PROD=PDGF associated protein /DB_XREF=gi:7657440 /UG=Hs.278426 PDGFA associated protein 1 /FL=gb:U41745.1 gb:BC000684.1 gb:NM_014891.1	NM_014891		NP_055706
202293_at	0.01714	Consensus includes gb:AW168948 /FEA=EST /DB_XREF=gi:6400473 /DB_XREF=est:xj15f07.x1 /CLONE=IMAGE:2657317 /UG=Hs.286148 stromal antigen 1 /FL=gb:NM_005862.1	NM_005862		NP_005853
202294_at	0.034296	Consensus includes gb:AI126490 /FEA=EST /DB_XREF=gi:3595004 /DB_XREF=est:qd82h04.x1 /CLONE=IMAGE:1736023 /UG=Hs.286148 stromal antigen 1 /FL=gb:NM_005862.1	NM_005862		NP_005853
202295_s_at	7.76E-04	gb:NM_004390.1 /DEF=Homo sapiens cathepsin H (CTSH), mRNA. /FEA=mRNA /GEN=CTSH /PROD=cathepsin H /DB_XREF=gi:4758095 /UG=Hs.288181 cathepsin H /FL=gb:BC002479.1 gb:NM_004390.1	NM_004390		NP_683880

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202297_s_at	0.001812	gb:AF157324.1 /DEF=Homo sapiens RER1 protein (RER1) mRNA, complete cds. /FEA=mRNA /GEN=RER1 /PROD=RER1 protein /DB_XREF=gi:7688698 /UG=Hs.40500 similar to S. cerevisiae RER1 /FL=gb:BC004965.1 gb:NM_007033.1 gb:AF157324.1	AF157324		NP_008964
202298_at	7.25E-04	gb:NM_004541.2 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1 (7.5kD, MWFE) (NDUFA1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=NDUFA1 /PROD=NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1 /DB_XREF=gi:13699820 /UG=Hs.74823 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1 (7.5kD, MWFE) /FL=gb:BC000266.1 gb:NM_004541.2 gb:U54993.1	NM_004541		NP_004532
202299_s_at	6.66E-04	gb:NM_006402.1 /DEF=Homo sapiens hepatitis B virus x-interacting protein (9.6kD) (XIP), mRNA. /FEA=mRNA /GEN=XIP /PROD=hepatitis B virus x-interacting protein /DB_XREF=gi:5454169 /UG=Hs.80464 hepatitis B virus x-interacting protein (9.6kD) /FL=gb:AF029890.1 gb:NM_006402.1	NM_006402		NP_006393

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202300_at	7.63E-04	gb:NM_006402.1 /DEF=Homo sapiens hepatitis B virus x-interacting protein (9.6kD) (XIP), mRNA. /FEA=mRNA /GEN=XIP. /PROD=hepatitis B virus x-interacting protein /DB_XREF=gi:5454169 /UG=Hs.80464 hepatitis B virus x-interacting protein (9.6kD) /FL=gb:AF029890.1 gb:NM_006402.1	NM_006402		NP_006393
202304_at	0.002724	gb:NM_014923.1 /DEF=Homo sapiens KIAA0970 protein (KIAA0970), mRNA. /FEA=mRNA /GEN=KIAA0970 /PROD=KIAA0970 protein /DB_XREF=gi:7662419 /UG=Hs.103329 KIAA0970 protein /FL=gb:AB023187.1 gb:NM_014923.1	NM_014923		
202313_at	0.00121	gb:NM_002717.1 /DEF=Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform (PPP2R2A), mRNA. /FEA=mRNA /GEN=PPP2R2A /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform /DB_XREF=gi:4506018 /UG=Hs.179574 protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform /FL=gb:M64929.1 gb:NM_002717.1	NM_002717		NP_002708

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
202317_s_at	0.047555	gb:NM_006048.1 /DEF=Homo sapiens ubiquitination factor E4B (homologous to yeast UFD2) (UBE4B), mRNA. /FEA=mRNA /GEN=UBE4B /PROD=ubiquitination factor E4B (homologous to yeast UFD2) /DB_XREF=gi:5174482 /UG=Hs.24594 ubiquitination factor E4B (homologous to yeast UFD2) /FL=gb:AF043117.1 gb:NM_006048.1	NM_006048		NP_006039
202325_s_at	0.003965	gb:NM_001685.1 /DEF=Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit F6 (ATP5J), mRNA. /FEA=mRNA /GEN=ATP5J /PROD=ATP synthase, H ⁺ transporting, mitochondrial /DB_XREF=gi:4502292 /UG=Hs.73851 ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit F6 /FL=gb:M37104.1 gb:M73031.1 gb:NM_001685.1 gb:AL110183.1	NM_001685		NP_001676
202329_at	0.018534	gb:NM_004383.1 /DEF=Homo sapiens c-src tyrosine kinase (CSK), mRNA. /FEA=mRNA /GEN=CSK /PROD=c-src tyrosine kinase /DB_XREF=gi:4758077 /UG=Hs.77793 c-src tyrosine kinase /FL=gb:NM_004383.1	NM_004383		NP_004374
202330_s_at	0.033831	gb:NM_003362.1 /DEF=Homo sapiens uracil-DNA glycosylase (UNG), mRNA. /FEA=mRNA /GEN=UNG /PROD=uracil-DNA glycosylase /DB_XREF=gi:6224978 /UG=Hs.78853 uracil-DNA glycosylase /FL=gb:NM_003362.1	NM_003362		NP_550433

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202333_s_at	0.011969	ubiquitin-conjugating enzyme E2B (RAD6 homolog)	AA877765	Hs.811	NP_003328
202334_s_at	0.005574	ubiquitin-conjugating enzyme E2B (RAD6 homolog)	AA877765	Hs.811	NP_003328
202343_x_at	0.001852	gb:NM_001862.1 /DEF=Homo sapiens cytochrome c oxidase subunit Vb (COX5B), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=COX5B /PROD=cytochrome c oxidase subunit Vb /DB_XREF=gi:4502982 /UG=Hs.1342 cytochrome c oxidase subunit Vb. /FL=gb:M19961.1 gb:NM_001862.1	NM_001862		NP_001853
202346_at	7.76E-04	gb:NM_005339.2 /DEF=Homo sapiens huntingtin interacting protein 2 (HIP2), mRNA. /FEA=mRNA /GEN=HIP2 /PROD=huntingtin interacting protein 2 /DB_XREF=gi:12545382 /UG=Hs.155485 huntingtin interacting protein 2 /FL=gb:NM_005339.2 gb:U58522.1 gb:AB022435.1	NM_005339		NP_005330
202347_s_at	0.002724	gb:AB022435.1 /DEF=Homo sapiens LIG mRNA for E2 ubiquitin-conjugating enzyme, complete cds. /FEA=mRNA /GEN=LIG /PROD=E2 ubiquitin conjugating enzyme /DB_XREF=gi:4996607 /UG=Hs.155485 huntingtin interacting protein 2 /FL=gb:NM_005339.2 gb:U58522.1 gb:AB022435.1	AB022435		NP_005330

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
202349_at	8.28E-04	gb:NM_000113.1 /DEF=Homo sapiens dystonia 1, torsion (autosomal dominant; torsin A) (DYT1), mRNA. /FEA=mRNA /GEN=DYT1 /PROD=torsin A /DB_XREF=gi:4557540 /UG=Hs.19261 dystonia 1, torsion (autosomal dominant; torsin A) /FL=gb:BC000674.1 gb:AF007871.1 gb:NM_000113.1	NM_000113		NP_000104
202352_s_at	7.63E-04	proteasome (prosome, macropain) 26S subunit, non-ATPase, 12	AI446530	Hs.4295	NP_777360
202358_s_at	0.015264	KIAA0254 gene product	BG434168	Hs.76906	NP_055573
202360_at	7.25E-04	gb:NM_014757.2 /DEF=Homo sapiens mastermind (Drosophila), homolog of (MAML1), mRNA. /FEA=mRNA /GEN=MAML1 /PROD=homolog of Drosophila mastermind /DB_XREF=gi:13376996 /UG=Hs.76986 mastermind (Drosophila), homolog of /FL=gb:NM_014757.2 gb:D83785.1	NM_014757		NP_055572
202362_at	0.003875	gb:NM_002884.1 /DEF=Homo sapiens RAP1A, member of RAS oncogene family (RAP1A), mRNA. /FEA=mRNA /GEN=RAP1A /PROD=RAP1A, member of RAS oncogene family /DB_XREF=gi:4506412 /UG=Hs.865 RAP1A, member of RAS oncogene family /FL=gb:NM_002884.1 gb:M22995.1	NM_002884		NP_002875

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
G n Id ntifier	p-value	Description	Gene Accession No.	Unigene Acc ssion No.	Protein Accession No.
202364_at	0.001812	gb:NM_005962.1 /DEF=Homo sapiens MAX-interacting protein 1 (MXI1), mRNA. /FEA=mRNA /GEN=MXI1 /PROD=MAX-interacting protein 1. /DB_XREF=gi:5174596 /UG=Hs.118630 MAX-interacting protein 1. /FL=gb:L07648.1 gb:NM_005962.1 gb:D63940.1	NM_005962		NP_569157
202367_at	0.00333	gb:NM_001913.1 /DEF=Homo sapiens cut (Drosophila)-like 1 (CCAAT displacement protein) (CUTL1), mRNA. /FEA=mRNA /GEN=CUTL1 /PROD=cut (Drosophila)-like 1 (CCAAT displacementprotein) /DB_XREF=gi:4503168 /UG=Hs.147049 cut (Drosophila)-like 1 (CCAAT displacement protein) /FL=gb:NM_001913.1 gb:L12579.1	NM_001913		NP_001904
202371_at	0.026914	gb:NM_024863.1 /DEF=Homo sapiens hypothetical protein FLJ21174 (FLJ21174), mRNA. /FEA=mRNA /GEN=FLJ21174 /PROD=hypothetical protein FLJ21174 /DB_XREF=gi:13376293 /UG=Hs.194329 hypothetical protein FLJ21174 /FL=gb:AF271783.1 gb:NM_024863.1	NM_024863		NP_079139
202373_s_at	0.007174	gb:AF255648.1 /DEF=Homo sapiens rGAP-iso mRNA, complete cds. /FEA=mRNA /PROD=rGAP-iso /DB_XREF=gi:12005820 /UG=Hs.197289 rab3 GTPase-activating protein, non-catalytic subunit (150kD) /FL=gb:AF255648.1 gb:AF004828.1 gb:NM_012414.1	AF255648		NP_036546

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202374_s_at	0.003595	gb:NM_012414.1 /DEF=Homo sapiens rab3 GTPase-activating protein, non-catalytic subunit (150kD) (RAB3-GAP150), mRNA. /FEA=mRNA /GEN=RAB3-GAP150 /PROD=rab3 GTPase-activating protein, non-catalytic subunit (150kD) /DB_XREF=gi:6912619 /UG=Hs.197289 rab3 GTPase-activating protein, non-catalytic subunit (150kD) /FL=gb:AF255648.1 gb:AF004828.1 gb:NM_012414.1	NM_012414		NP_036546
202375_at	7.76E-04	gb:NM_014822.1 /DEF=Homo sapiens SEC24 (S. cerevisiae) related gene family, member D (SEC24D), mRNA. /FEA=mRNA /GEN=SEC24D /PROD=SEC24 (S. cerevisiae) related gene family, member D /DB_XREF=gi:7662658 /UG=Hs.19822 SEC24 (S. cerevisiae) related gene family, member D /FL=gb:AB018298.1 gb:AF130464.2 gb:NM_014822.1	NM_014822		NP_055637
202377_at	6.66E-04	ESTs	AW026535	Hs.23581	NP_059996
202379_s_at	6.66E-04	natural killer-tumor recognition sequence	AI361805	Hs.241493	NP_005376

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202381_at	6.66E-04	gb:NM_003816.1 /DEF=Homo sapiens a disintegrin and metalloproteinase domain 9 (meltrin gamma) (ADAM9), mRNA. /FEA=mRNA /GEN=ADAM9 /PROD=a disintegrin and metalloproteinase domain 9preproprotein /DB_XREF=gi:4501914 /UG=Hs.2442 a disintegrin and metalloproteinase domain 9 (meltrin gamma) /FL=gb:U41766.1 gb:NM_003816.1	NM_003816		NP_003807
202382_s_at	0.00453	gb:NM_005471.1 /DEF=Homo sapiens glucosamine-6-phosphate isomerase (GNPI), mRNA. /FEA=mRNA /GEN=GNPI /PROD=glucosamine-6-phosphate isomerase /DB_XREF=gi:13027377 /UG=Hs.278500 glucosamine-6-phosphate isomerase /FL=gb:NM_005471.1 gb:AF029914.1 gb:AF048826.1 gb:D31766.1	NM_005471		NP_005462
202386_s_at	7.25E-04	gb:NM_019081.1 /DEF=Homo sapiens KIAA0430 gene product (KIAA0430), mRNA. /FEA=mRNA /GEN=KIAA0430 /PROD=hypothetical protein A-362G6.1 /DB_XREF=gi:11464998 /UG=Hs.30909 KIAA0430 gene product /FL=gb:NM_019081.1	NM_019081		NP_061954

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
202387_at	7.63E-04	gb:NM_004323.2 /DEF=Homo sapiens BCL2-associated athanogene (BAG1), mRNA. /FEA=mRNA /GEN=BAG1 /PROD=BCL2-associated athanogene /DB_XREF=gi:7549801 /UG=Hs.41714 BCL2-associated athanogene /FL=gb:BC001936.1 gb:AF022224.1 gb:U46917.1 gb:NM_004323.2	NM_004323		NP_004314
202391_at	0.024598	gb:NM_006317.1 /DEF=Homo sapiens brain acid-soluble protein 1 (BASP1), mRNA. /FEA=mRNA /GEN=BASP1 /PROD=brain acid-soluble protein 1 /DB_XREF=gi:5453749 /UG=Hs.79516 brain abundant, membrane attached signal protein 1 /FL=gb:BC000518.1 gb:AF039656.1 gb:NM_006317.1	NM_006317		NP_006308
202393_s_at	0.002603	gb:NM_005655.1 /DEF=Homo sapiens TGFB inducible early growth response (TIEG), mRNA. /FEA=mRNA /GEN=TIEG /PROD=TGFB inducible early growth response /DB_XREF=gi:5032176 /UG=Hs.82173 TGFB inducible early growth response /FL=gb:U21847.1 gb:NM_005655.1	NM_005655		NP_005646

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202395_at	7.76E-04	gb:NM_006178.1 /DEF=Homo sapiens N-ethylmaleimide-sensitive factor (NSF), mRNA. /FEA=mRNA /GEN=NSF /PROD=N-ethylmaleimide-sensitive factor /DB_XREF=gi:11079227 /UG=Hs.108802 N-ethylmaleimide-sensitive factor /FL=gb:AF102846.2 gb:NM_006178.1 gb:AF135168.1	NM_006178		NP_006169
202397_at	0.014602	gb:NM_005796.1 /DEF=Homo sapiens nuclear transport factor 2 (placental protein 15) (PP15), mRNA. /FEA=mRNA /GEN=PP15 /PROD=nuclear transport factor 2 (placental protein15) /DB_XREF=gi:5031984 /UG=Hs.151734 nuclear transport factor 2 (placental protein 15) /FL=gb:U43939.1 gb:BC002348.1 gb:NM_005796.1	NM_005796		NP_005787
202398_at	0.028996	gb:BC002785.1 /DEF=Homo sapiens, adaptor-related protein complex 3, sigma 2 subunit, clone MGC:3577, mRNA, complete cds. /FEA=mRNA /PROD=adaptor-related protein complex 3, sigma 2subunit /DB_XREF=gi:12803880 /UG=Hs.154782 adaptor-related protein complex 3, sigma 2 subunit /FL=gb:BC002785.1 gb:NM_005829.1	BC002785		NP_005820

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202399_s_at	7.63E-04	gb:NM_005829.1 /DEF=Homo sapiens adaptor-related protein complex 3, sigma 2 subunit (AP3S2), mRNA. /FEA=mRNA /GEN=AP3S2 /PROD=adaptor-related protein complex 3, sigma 2 subunit /DB_XREF=gi:5031580 /UG=Hs.154782 adaptor-related protein complex 3, sigma 2 subunit /FL=gb:BC002785.1 gb:NM_005829.1	NM_005829		NP_005820
202401_s_at	0.004249	gb:NM_003131.1 /DEF=Homo sapiens serum response factor (c-fos serum response element-binding transcription factor) (SRF), mRNA. /FEA=mRNA /GEN=SRF /PROD=serum response factor (c-fos serum response element-binding transcription factor) /DB_XREF=gi:4507204 /UG=Hs.155321 serum response factor (c-fos serum response element-binding transcription factor) /FL=gb:J03161.1 gb:NM_003131.1	NM_003131		NP_003122
202402_s_at	0.01063	gb:NM_001751.1 /DEF=Homo sapiens cysteinyl-tRNA synthetase (CARS), mRNA. /FEA=mRNA /GEN=CARS /PROD=cysteinyl-tRNA synthetase /DB_XREF=gi:10835050 /UG=Hs.159604 cysteinyl-tRNA synthetase /FL=gb:NM_001751.1 gb:BC002880.1 gb:AF288206.1 gb:AF288207.1	NM_001751		NP_644802
202412_s_at	0.028771	ubiquitin specific protease 1	AW499935	Hs.35086	NP_003359

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
202422_s_at	0.009486	gb:NM_022977.1 /DEF=Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 4 (FACL4), transcript variant 2, mRNA. /FEA=mRNA /GEN=FACL4 /PROD=long-chain fatty-acid-Coenzyme A ligase 4, isoform 2 /DB_XREF=gi:12669908 /UG=Hs.81452 fatty-acid-Coenzyme A ligase, long-chain 4 /FL=gb:NM_022977.1	NM_022977		NP_075266
202423_at	0.013199	gb:NM_006766.1 /DEF=Homo sapiens zinc finger protein 220 (ZNF220), mRNA. /FEA=mRNA /GEN=ZNF220 /PROD=zinc finger protein 220 /DB_XREF=gi:5803097 /UG=Hs.82210 zinc finger protein 220 /FL=gb:U47742.1 gb:NM_006766.1	NM_006766		NP_006757
202427_s_at	0.001479	gb:NM_015415.1 /DEF=Homo sapiens DKFZP564B167 protein (DKFZP564B167), mRNA. /FEA=mRNA /GEN=DKFZP564B167 /PROD=DKFZP564B167 protein /DB_XREF=gi:7661601 /UG=Hs.76285 DKFZP564B167 protein /FL=gb:AL110297.1 gb:NM_015415.1	NM_015415		NP_056230
202429_s_at	0.009378	gb:AL353950.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761L0516 (from clone DKFZp761L0516); complete cds. /FEA=mRNA /GEN=DKFZp761L0516 /PROD=hypothetical protein /DB_XREF=gi:7669991 /UG=Hs.272458 protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha) /FL=gb:J05480.1 gb:L14778.1 gb:NM_000944.1 gb:AL353950.1	AL353950		NP_000935

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202430_s_at	0.002446	gb:NM_021105.1 /DEF=Homo sapiens phospholipid scramblase 1 (PLSCR1), mRNA. /FEA=mRNA /GEN=PLSCR1 /PROD=phospholipid scramblase 1 /DB_XREF=gi:10863876 /UG=Hs.198282 phospholipid scramblase 1 /FL=gb:NM_021105.1 gb:AB006746.1 gb:AF098642.1	NM_021105		NP_066928
202432_at	0.001406	gb:NM_021132.1 /DEF=Homo sapiens protein phosphatase 3 (formerly 2B), catalytic subunit, beta isoform (calcineurin A beta) (PPP3CB), mRNA. /FEA=mRNA /GEN=PPP3CB /PROD=protein phosphatase 3 (formerly 2B), catalytic subunit, beta isoform (calcineurin A beta) /DB_XREF=gi:11036639 /UG=Hs.151531 protein phosphatase 3 (formerly 2B), catalytic subunit, beta isoform (calcineurin A beta) /FL=gb:NM_021132.1 gb:M29551.1	NM_021132		NP_066955
202433_at	7.63E-04	gb:NM_005827.1 /DEF=Homo sapiens UDP-galactose transporter related (UGTREL1), mRNA. /FEA=mRNA /GEN=UGTREL1 /PROD=UDP-galactose transporter related /DB_XREF=gi:5032212 /UG=Hs.154073 UDP-galactose transporter related /FL=gb:D87989.1 gb:NM_005827.1	NM_005827		NP_005818

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202435_s_at	0.039362	Consensus includes gb:AU154504 /FEA=EST /DB_XREF=gi:11016025 /DB_XREF=est:AU154504 /CLONE=NT2RP4001328 /UG=Hs.154654 cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile) /FL=gb:NM_000104.2 gb:U03688.1	NM_000104		NP_000095
202436_s_at	0.049392	Consensus includes gb:AU144855 /FEA=EST /DB_XREF=gi:11006376 /DB_XREF=est:AU144855 /CLONE=HEMBA1003161 /UG=Hs.154654 cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile) /FL=gb:NM_000104.2 gb:U03688.1	NM_000104		NP_000095
202437_s_at	0.008944	gb:NM_000104.2 /DEF=Homo sapiens cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile) (CYP1B1), mRNA /FEA=mRNA /GEN=CYP1B1 /PROD=cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 /DB_XREF=gi:13325059 /UG=Hs.154654 cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile) /FL=gb:NM_000104.2 gb:U03688.1	NM_000104		NP_000095
202441_at	7.63E-04	similar to Caenorhabditis elegans protein C42C1.9	AL568449	Hs.285818	NP_006450

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202442_at	7.25E-04	gb:NM_001284.1 /DEF=Homo sapiens adaptor-related protein complex 3, sigma 1 subunit (AP3S1), mRNA. /FEA=mRNA /GEN=AP3S1 /PROD=adaptor-related protein complex 3, sigma 1 subunit /DB_XREF=gi:4502860 /UG=Hs.80917 adaptor-related protein complex 3, sigma 1 subunit /FL=gb:BC000804.1 gb:D63643.1 gb:U91932.1 gb:NM_001284.1	NM_001284		NP_001275
202443_x_at	0.002603	Notch homolog 2 (Drosophila)	AA291203	Hs.131286	NP_077719
202445_s_at	0.033358	gb:NM_024408.1 /DEF=Homo sapiens Notch (Drosophila) homolog 2 (NOTCH2), mRNA. /FEA=mRNA /GEN=NOTCH2 /PROD=Notch (Drosophila) homolog 2. /DB_XREF=gi:13249343 /UG=Hs.8121 Notch (Drosophila) homolog 2 /FL=gb:AF315356.1 gb:NM_024408.1	NM_024408		NP_077719
202446_s_at	0.003669	phospholipid scramblase 1	AI825926	Hs.198282	NP_066928
202447_at	0.006596	gb:NM_001359.1 /DEF=Homo sapiens 2,4-dienoyl CoA reductase 1, mitochondrial (DECR1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=DECR1 /PROD=2,4-dienoyl CoA reductase 1 precursor /DB_XREF=gi:4503300 /UG=Hs.81548 2,4-dienoyl CoA reductase 1, mitochondrial /FL=gb:U49352.1 gb:NM_001359.1 gb:L26050.1	NM_001359		NP_001350
202457_s_at	0.003976	protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha)	AA911231	Hs.272458	NP_000935

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
202462_s_at	0.007839	gb:NM_014829.1 /DEF=Homo sapiens KIAA0801 gene product (KIAA0801), mRNA. /FEA=mRNA /GEN=KIAA0801 /PROD=KIAA0801 gene product /DB_XREF=gi:7662317 /UG=Hs.17585 KIAA0801 gene product /FL=gb:AB018344.1 gb:NM_014829.1	NM_014829		NP_055644
202464_s_at	0.00121	gb:NM_004566.1 /DEF=Homo sapiens 6-phosphofructo-2-kinasefructose-2,6-biphosphatase 3 (PFKFB3), mRNA. /FEA=mRNA /GEN=PFKFB3 /PROD=6-phosphofructo-2-kinasefructose-2,6-biphosphatase 3 /DB_XREF=gi:4758899 /UG=Hs.195471 6-phosphofructo-2-kinasefructose-2,6-biphosphatase 3 /FL=gb:D49817.1 gb:AF109735.1 gb:NM_004566.1	NM_004566		NP_004557
202469_s_at	0.003965	cleavage and polyadenylation specific factor 6, 68kDa	AU149367	Hs.64542	
202471_s_at	0.013527	gb:NM_004135.1 /DEF=Homo sapiens isocitrate dehydrogenase 3 (NAD+) gamma (IDH3G), mRNA. /FEA=mRNA /GEN=IDH3G /PROD=isocitrate dehydrogenase 3 (NAD+) gamma /DB_XREF=gi:4758581 /UG=Hs.75253 isocitrate dehydrogenase 3 (NAD+) gamma /FL=gb:BC001902.1 gb:BC000933.2 gb:U40272.1 gb:NM_004135.1	NM_004135		NP_777358

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Acc ssion No.
202474_s_at	0.018839	gb:NM_005334.1 /DEF=Homo sapiens host cell factor C1 (VP16-accessory protein) (HCFC1), mRNA. /FEA=mRNA /GEN=HCFC1 /PROD=host cell factor C1 (VP16-accessory protein) /DB_XREF=gi:4885402 /UG=Hs.83634 host cell factor C1 (VP16-accessory protein) /FL=gb:NM_005334.1	NM_005334		NP_005325
202478_at	0.004519	gb:NM_021643.1 /DEF=Homo sapiens GS3955 protein (GS3955), mRNA. /FEA=mRNA /GEN=GS3955 /PROD=GS3955 protein /DB_XREF=gi:11056053 /UG=Hs.155418 GS3955 protein /FL=gb:NM_021643.1 gb:BC002637.1 gb:D87119.1	NM_021643		NP_067675
202479_s_at	0.049223	gb:BC002637.1 /DEF=Homo sapiens, GS3955 protein, clone MGC:3860, mRNA, complete cds. /FEA=mRNA /PROD=GS3955 protein /DB_XREF=gi:12803604 /UG=Hs.155418 GS3955 protein /FL=gb:NM_021643.1 gb:BC002637.1 gb:D87119.1	BC002637		NP_067675
202480_s_at	0.003366	gb:NM_004216.1 /DEF=Homo sapiens death effector domain-containing (DEDD), mRNA. /FEA=mRNA /GEN=DEDD /PROD=death effector domain-containing /DB_XREF=gi:4758143 /UG=Hs.169681 death effector domain-containing /FL=gb:AF083236.1 gb:AF043733.1 gb:AF100341.1 gb:NM_004216.1	NM_004216		NP_127491

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
202484_s_at	0.005652	gb:AF072242.1 /DEF=Homo sapiens methyl-CpG binding protein MBD2 (MBD2) mRNA, complete cds. /FEA=mRNA /GEN=MBD2 /PROD=methyl-CpG binding protein MBD2 /DB_XREF=gi:3800792 /UG=Hs.25674 methyl-CpG binding domain protein 2 /FL=gb:AF072242.1 gb:NM_003927.2	AF072242		NP_056647
202485_s_at	0.04323	gb:NM_003927.2 /DEF=Homo sapiens methyl-CpG binding domain protein 2 (MBD2), transcript variant 1, mRNA. /FEA=mRNA /GEN=MBD2 /PROD=methyl-CpG binding domain protein 2, isoform 1 /DB_XREF=gi:7710146 /UG=Hs.25674 methyl-CpG binding domain protein 2 /FL=gb:AF072242.1 gb:NM_003927.2	NM_003927		NP_056647
202487_s_at	6.66E-04	gb:NM_012412.1 /DEF=Homo sapiens purine-rich element binding protein B (PURB), mRNA. /FEA=mRNA /GEN=PURB /PROD=purine-rich element binding protein B /DB_XREF=gi:6912615 /UG=Hs.301005 purine-rich element binding protein B. /FL=gb:BC000098.1 gb:BC004274.1 gb:AF081192.1 gb:NM_012412.1	NM_012412		NP_619541
202492_at	0.039831	gb:NM_024085.1 /DEF=Homo sapiens hypothetical protein FLJ22169 (FLJ22169), mRNA. /FEA=mRNA /GEN=FLJ22169 /PROD=hypothetical protein FLJ22169 /DB_XREF=gi:13129081 /UG=Hs.323363 hypothetical protein FLJ22169 /FL=gb:BC001206.1 gb:NM_024085.1	NM_024085		NP_076990

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202495_at	7.76E-04	gb:NM_003192.1 /DEF=Homo sapiens tubulin-specific chaperone c (TBCC), mRNA. /FEA=mRNA /GEN=TBCC /PROD=beta-tubulin cofactor C /DB_XREF=gi:4507372 /UG=Hs.75064 tubulin-specific chaperone c /FL=gb:U61234.1 gb:NM_003192.1	NM_003192		NP_003183
202499_s_at	9.85E-04	gb:NM_006931.1 /DEF=Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 3 (SLC2A3), mRNA. /FEA=mRNA /GEN=SLC2A3 /PROD=solute carrier family 2 (facilitated glucosetransporter), member 3 /DB_XREF=gi:5902089 /UG=Hs.7594 solute carrier family 2 (facilitated glucose transporter), member 3 /FL=gb:M20681.1 gb:NM_006931.1	NM_006931		NP_008862
202503_s_at	7.25E-04	gb:NM_014736.1 /DEF=Homo sapiens KIAA0101 gene product (KIAA0101), mRNA. /FEA=mRNA /GEN=KIAA0101 /PROD=KIAA0101 gene product /DB_XREF=gi:7661905 /UG=Hs.81892 KIAA0101 gene product /FL=gb:D14657.1 gb:NM_014736.1	NM_014736		NP_055551
202506_at	0.002724	gb:NM_006751.1 /DEF=Homo sapiens sperm specific antigen 2 (SSFA2), mRNA. /FEA=mRNA /GEN=SSFA2 /PROD=sperm specific antigen 2 /DB_XREF=gi:5803178 /UG=Hs.82767 sperm specific antigen 2 /FL=gb:M61199.1 gb:NM_006751.1	NM_006751		NP_006742

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202511_s_at	0.001167	Consensus includes gb:AK001899.1 /DEF=Homo sapiens cDNA FLJ11037 fis, clone PLACE1004316, highly similar to H.sapiens mRNA for apoptosis specific protein. /FEA=mRNA /DB_XREF=gi:7023451 /UG=Hs.11171 APG5 (autophagy 5, S. cerevisiae)-like /FL=gb:NM_004849.1	AK001899		NP_004840
202513_s_at	0.014602	gb:NM_006245.1 /DEF=Homo sapiens protein phosphatase 2, regulatory subunit B (B56), delta isoform (PPP2R5D), mRNA. /FEA=mRNA /GEN=PPP2R5D /PROD=protein phosphatase 2, regulatory subunit B(B56), delta isoform /DB_XREF=gi:5453953 /UG=Hs.118244 protein phosphatase 2, regulatory subunit B (B56), delta isoform /FL=gb:BC001095.1 gb:L76702.1 gb:AB000634.1 gb:NM_006245.1	NM_006245		NP_851308
202519_at	0.008874	gb:NM_014938.1 /DEF=Homo sapiens KIAA0867 protein (MONDOA), mRNA. /FEA=mRNA /GEN=MONDOA /PROD=MondoA protein /DB_XREF=gi:7662347 /UG=Hs.52081 KIAA0867 protein /FL=gb:AB020674.1 gb:NM_014938.1	NM_014938		NP_055753

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202521_at	0.011487	gb:NM_006565.1 /DEF=Homo sapiens CCCTC-binding factor (zinc finger protein) (CTCF), mRNA. /FEA=mRNA /GEN=CTCF /PROD=CCCTC-binding factor (zinc finger protein) /DB_XREF=gi:5729789 /UG=Hs.57419 CCCTC-binding factor (zinc finger protein) /FL=gb:NM_006565.1 gb:U25435.1	NM_006565		NP_006556
202524_s_at	0.040104	gb:NM_014767.1 /DEF=Homo sapiens KIAA0275 gene product (KIAA0275), mRNA. /FEA=mRNA /GEN=KIAA0275 /PROD=KIAA0275 gene product /DB_XREF=gi:7662035 /UG=Hs.74583 KIAA0275 gene product /FL=gb:D87465.1 gb:NM_014767.1	NM_014767		NP_055582
202527_s_at	0.009678	gb:NM_005359.1 /DEF=Homo sapiens MAD (mothers against decapentaplegic, Drosophila) homolog 4 (MADH4), mRNA. /FEA=mRNA /GEN=MADH4 /PROD=MAD (mothers against decapentaplegic, Drosophila) homolog 4 /DB_XREF=gi:4885456 /UG=Hs.75862 MAD (mothers against decapentaplegic, Drosophila) homolog 4 /FL=gb:U44378.1 gb:BC002379.1 gb:NM_005359.1	NM_005359		NP_005350

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202529_at	0.002109	gb:NM_002766.1 /DEF=Homo sapiens phosphoribosyl pyrophosphate synthetase-associated protein 1 (PRPSAP1), mRNA. /FEA=mRNA /GEN=PRPSAP1 /PROD=phosphoribosyl pyrophosphatesynthetase-associated protein 1 /DB_XREF=gi:4506130 /UG=Hs.77498 phosphoribosyl pyrophosphate synthetase-associated protein 1 /FL=gb:D61391.1 gb:NM_002766.1	NM_002766		NP_002757
202530_at	7.25E-04	gb:NM_001315.1 /DEF=Homo sapiens mitogen-activated protein kinase 14 (MAPK14), mRNA. /FEA=mRNA /GEN=MAPK14 /PROD=mitogen-activated protein kinase 14 /DB_XREF=gi:4503068 /UG=Hs.79107 mitogen-activated protein kinase 14 /FL=gb:NM_001315.1 gb:L35263.1	NM_001315		NP_620583
202532_s_at	0.005574	gb:BC000192.1 /DEF=Homo sapiens, dihydrofolate reductase, clone MGC:857; mRNA, complete cds. /FEA=mRNA /PROD=dihydrofolate reductase /DB_XREF=gi:12652874 /UG=Hs.83765 dihydrofolate reductase /FL=gb:BC000192.1 gb:BC003584.1 gb:NM_000791.2	BC000192		NP_000782

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202534_x_at	0.001835	gb:NM_000791.2 /DEF=Homo sapiens dihydrofolate reductase (DHFR), mRNA. /FEA=mRNA /GEN=DHFR /PROD=dihydrofolate reductase /DB_XREF=gi:7262376 /UG=Hs.83765 dihydrofolate reductase /FL=gb:BC000192.1 gb:BC003584.1 gb:NM_000791.2	NM_000791		NP_000782
202535_at	0.001008	gb:NM_003824.1 /DEF=Homo sapiens Fas (TNFRSF6)-associated via death domain (FADD), mRNA. /FEA=mRNA /GEN=FADD /PROD=Fas (TNFRSF6)-associated via death domain /DB_XREF=gi:4505228 /UG=Hs.86131 Fas (TNFRSF6)-associated via death domain /FL=gb:BC000334.1 gb:NM_003824.1 gb:U24231.1	NM_003824		NP_003815
202536_at	7.25E-04	Consensus includes gb:AK002165.1 /DEF=Homo sapiens cDNA FLJ11303 fis, clone PLACE1009995, highly similar to Homo sapiens mRNA; cDNA DKFZp564O123. /FEA=mRNA /DB_XREF=gi:7023876 /UG=Hs.11449 DKFZP564O123 protein /FL=gb:AF151842.1 gb:AL080122.1 gb:NM_014043.1	AK002165		NP_054762

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202538_s_at	7.63E-04	gb:NM_014043.1 /DEF=Homo sapiens DKFZP564O123 protein (DKFZP564O123), mRNA. /FEA=mRNA /GEN=DKFZP564O123 /PROD=DKFZP564O123 protein /DB_XREF=gi:7661633 /UG=Hs.11449 DKFZP564O123 protein /FL=gb:AF151842.1 gb:AL080122.1 gb:NM_014043.1	NM_014043		NP_054762
202539_s_at	7.25E-04	3-hydroxy-3-methylglutaryl-Coenzyme A reductase	AL518627	Hs.11899	NP_000850
202540_s_at	0.005652	gb:NM_000859.1 /DEF=Homo sapiens 3-hydroxy-3-methylglutaryl-Coenzyme A reductase (HMGCR), mRNA. /FEA=mRNA /GEN=HMGCR /PROD=3-hydroxy-3-methylglutaryl-Coenzyme A reductase /DB_XREF=gi:4557642 /UG=Hs.11899 3-hydroxy-3-methylglutaryl-Coenzyme A reductase. /FL=gb:M11058.1 gb:NM_000859.1	NM_000859		NP_000850
202543_s_at	0.003842	gb:BC005359.1 /DEF=Homo sapiens, glia maturation factor, beta, clone MGC:12462, mRNA, complete cds. /FEA=mRNA /PROD=glia maturation factor, beta /DB_XREF=gi:13529184 /UG=Hs.151413 glia maturation factor, beta /FL=gb:BC005359.1 gb:M86492.1 gb:AB001106.1 gb:NM_004124.1	BC005359		NP_004115

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
202544_at	0.023543	gb:NM_004124.1 /DEF=Homo sapiens glia maturation factor, beta (GMFB), mRNA. /FEA=mRNA /GEN=GMFB /PROD=glia maturation factor, beta /DB_XREF=gi:4758441 /UG=Hs.151413 glia maturation factor, beta /FL=gb:BC005359.1 gb:M86492.1 gb:AB001106.1 gb:NM_004124.1	NM_004124		NP_004115
202550_s_at	7.63E-04	gb:NM_004738.1 /DEF=Homo sapiens VAMP (vesicle-associated membrane protein)-associated protein B and C (VAPB), mRNA. /FEA=mRNA /GEN=VAPB /PROD=VAMP (vesicle-associated membraneprotein)-associated protein B and C /DB_XREF=gi:4759301 /UG=Hs.182625 VAMP (vesicle-associated membrane protein)-associated protein B and C /FL=gb:BC001712.1 gb:AF086628.1 gb:NM_004738.1 gb:AF160212.1	NM_004738		NP_004729
202553_s_at	0.005652	gb:NM_015484.1 /DEF=Homo sapiens GCIP-interacting protein p29 (P29), mRNA. /FEA=mRNA /GEN=P29 /PROD=GCIP-interacting protein p29 /DB_XREF=gi:7661635 /UG=Hs.20013 GCIP-interacting protein p29 /FL=gb:AF273089.1 gb:AL080166.1 gb:NM_015484.1	NM_015484		NP_056299

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniProt Accession No.	Prot in Accession No.
202556_s_at	0.044397	gb:NM_006337.1 /DEF=Homo sapiens microspherule protein 1 (MCRS1), mRNA. /FEA=mRNA /GEN=MCRS1 /PROD=microspherule protein 1 /DB_XREF=gi:5453693 /UG=Hs.25313 microspherule protein 1 /FL=gb:AF068007.1 gb:NM_006337.1	NM_006337		NP_006328
202565_s_at	6.66E-04	gb:NM_003174.2 /DEF=Homo sapiens supervillin (SVIL), transcript variant 1, mRNA. /FEA=mRNA /GEN=SVIL /PROD=supervillin, isoform 1 /DB_XREF=gi:11496980 /UG=Hs.154567 supervillin /FL=gb:NM_003174.2 gb:AF051850.1 gb:AF051851.1	NM_003174		NP_068506
202566_s_at	0.049173	gb:AF051851.1 /DEF=Homo sapiens supervillin mRNA, complete cds. /FEA=mRNA /PROD=supervillin /DB_XREF=gi:2961251 /UG=Hs.154567 supervillin /FL=gb:NM_003174.2 gb:AF051850.1 gb:AF051851.1	AF051851		NP_068506
202567_at	0.036998	gb:NM_004175.1 /DEF=Homo sapiens small nuclear ribonucleoprotein D3 polypeptide (18kD) (SNRPD3), mRNA. /FEA=mRNA /GEN=SNRPD3 /PROD=small nuclear ribonucleoprotein D3 polypeptide(18kD) /DB_XREF=gi:4759159 /UG=Hs.1575 small nuclear ribonucleoprotein D3 polypeptide (18kD) /FL=gb:BC000457.1 gb:BC003150.1 gb:NM_004175.1 gb:U15009.1	NM_004175		NP_004166
202568_s_at	0.002692	MAP/microtubule affinity-regulating kinase 3	AI745639	Hs.172766	NP_002367

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202569_s_at	0.007271	gb:NM_002376.1 /DEF=Homo sapiens MAPmicrotubule affinity-regulating kinase 3 (MARK3), mRNA. /FEA=mRNA /GEN=MARK3 /PROD=MAPmicrotubule affinity-regulating kinase 3 /DB_XREF=gi:4505102 /UG=Hs.172766 MAPmicrotubule affinity-regulating kinase 3 /FL=gb:M80359.1 gb:NM_002376.1	NM_002376		NP_002367
202582_s_at	7.63E-04	gb:AF306510.1 /DEF=Homo sapiens RANBPM mRNA, complete cds. /FEA=mRNA /PROD=RANBPM /DB_XREF=gi:13194575 /UG=Hs.279886 RAN binding protein 9 /FL=gb:AF306510.1 gb:AB008515.1 gb:NM_005493.1	AF306510		NP_005484
202583_s_at	7.76E-04	gb:NM_005493.1 /DEF=Homo sapiens RAN binding protein 9 (RANBP9), mRNA. /FEA=mRNA /GEN=RANBP9 /PROD=RAN binding protein 9 /DB_XREF=gi:4885570 /UG=Hs.279886 RAN binding protein 9 /FL=gb:AF306510.1 gb:AB008515.1 gb:NM_005493.1	NM_005493		NP_005484
202589_at	6.73E-04	gb:NM_001071.1 /DEF=Homo sapiens thymidylate synthetase (TYMS), mRNA. /FEA=mRNA /GEN=TYMS /PROD=thymidylate synthetase /DB_XREF=gi:4507750 /UG=Hs.82962 thymidylate synthetase /FL=gb:BC002567.1 gb:NM_001071.1	NM_001071		NP_001062

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202592_at	0.007818	gb:NM_001487.1 /DEF=Homo sapiens GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 1 (GCN5L1), mRNA. /FEA=mRNA /GEN=GCN5L1 /PROD=GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 1 /DB_XREF=gi:4503954 /UG=Hs.94672 GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 1 /FL=gb:D64007.1 gb:NM_001487.1	NM_001487		NP_001478
202596_at	0.001179	gb:BC000436.1 /DEF=Homo sapiens, endosulfine alpha, clone MGC:8394, mRNA, complete cds. /FEA=mRNA /PROD=endosulfine alpha /DB_XREF=gi:12653334 /UG=Hs.111680 endosulfine alpha /FL=gb:BC000436.1 gb:BC004461.1 gb:NM_004436.1 gb:AF157509.1	BC000436		NP_004427
202600_s_at	0.034999	nuclear receptor interacting protein 1	AI824012	Hs.155017	NP_003480
202603_at	0.001406	a disintegrin and metalloproteinase domain 10,	N51370	Hs.172028	NP_001101
202605_at	0.027672	gb:NM_000181.1 /DEF=Homo sapiens glucuronidase, beta (GUSB), mRNA. /FEA=mRNA /GEN=GUSB /PROD=glucuronidase, beta /DB_XREF=gi:4504222 /UG=Hs.183868 glucuronidase, beta /FL=gb:M15182.1 gb:NM_000181.1	NM_000181		NP_000172

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202606_s_at	0.024403	gb:NM_012290.1 /DEF=Homo sapiens tousled-like kinase 1 (TLK1), mRNA. /FEA=mRNA /GEN=TLK1 /PROD=tousled-like kinase 1 /DB_XREF=gi:6912719 /UG=Hs.18895 tousled-like kinase 1 /FL=gb:AB004885.1 gb:NM_012290.1 gb:AF246219.1	NM_012290		NP_036422
202607_at	0.011056	MAD2 mitotic arrest deficient-like 1 (yeast)	AL526632	Hs.79078	NP_001534
202610_s_at	7.76E-04	gb:AF135802.1 /DEF=Homo sapiens thyroid hormone receptor-associated protein complex component TRAP170 mRNA, complete cds. /FEA=mRNA /PROD=thyroid hormone receptor-associated protein complex component TRAP170 /DB_XREF=gi:4580325 /UG=Hs.21586 cofactor required for Sp1 transcriptional activation, subunit 2 (150kD) /FL=gb:AF304448.1 gb:AB006651.1 gb:AF104256.1 gb:AF135802.1 gb:NM_004229.1	AF135802		NP_004220
202611_s_at	0.003965	cofactor required for Sp1 transcriptional activation, subunit 2, 150kDa	AI971089	Hs.21586	NP_004220
202615_at	0.00643	guanine nucleotide binding protein (G protein), q polypeptide	BF222895	Hs.380144	
202621_at	7.25E-04	gb:NM_001571.1 /DEF=Homo sapiens interferon regulatory factor 3 (IRF3), mRNA. /FEA=mRNA /GEN=IRF3 /PROD=interferon regulatory factor 3 /DB_XREF=gi:4504724 /UG=Hs.75254 interferon regulatory factor 3 /FL=gb:NM_001571.1	NM_001571		NP_001562

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
202623_at	0.030776	gb:NM_018453.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM036 (BM036), mRNA. /FEA=mRNA /GEN=BM036 /PROD=uncharacterized bone marrow protein BM036 /DB_XREF=gi:8922092 /UG=Hs.7731 uncharacterized bone marrow protein BM036 /FL=gb:BC001245.1 gb:AF217512.1 gb:NM_018453.1	NM_018453		NP_060923
202625_at	0.014961	v-yes-1 Yamaguchi sarcoma viral related oncogene homolog	A1356412	Hs.80887	NP_002341
202629_at	0.004987	amyloid beta precursor protein (cytoplasmic tail) binding protein 2	AA046411	Hs.84084	NP_006371
202633_at	0.043116	gb:NM_007027.1 /DEF=Homo sapiens topoisomerase (DNA) II binding protein (TOPBP1), mRNA. /FEA=mRNA /GEN=TOPBP1 /PROD=topoisomerase (DNA) II binding protein /DB_XREF=gi:5902137 /UG=Hs.91417 topoisomerase (DNA) II binding protein /FL=gb:AB019397.1 gb:NM_007027.1	NM_007027		NP_008958
202635_s_at	0.004519	gb:NM_005034.1 /DEF=Homo sapiens polymerase (RNA) II (DNA directed) polypeptide K (7.0kD) (POLR2K), mRNA. /FEA=mRNA /GEN=POLR2K /PROD=polymerase (RNA) II (DNA directed) polypeptide K(7.0kD) /DB_XREF=gi:4826923 /UG=Hs.150675 polymerase (RNA) II (DNA directed) polypeptide K (7.0kD) /FL=gb:BC000806.1 gb:NM_005034.1	NM_005034		NP_005025

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
202636_at	0.002109	gb:NM_005667.1 /DEF=Homo sapiens zinc finger protein homologous to Zfp103 in mouse (ZFP103), mRNA. /FEA=mRNA /GEN=ZFP103 /PROD=zinc finger protein homologous to Zfp103 in mouse /DB_XREF=gi:5031824 /UG=Hs.155968 zinc finger protein homologous to Zfp103 in mouse /FL=gb:D76444.1 gb:NM_005667.1	NM_005667		NP_005658
202637_s_at	0.034925	intercellular adhesion molecule 1 (CD54), human rhinovirus receptor	AI608725	Hs.168383	NP_000192
202638_s_at	0.023543	gb:NM_000201.1 /DEF=Homo sapiens intercellular adhesion molecule 1 (CD54), human rhinovirus receptor (ICAM1), mRNA. /FEA=mRNA /GEN=ICAM1 /PROD=intercellular adhesion molecule 1 precursor /DB_XREF=gi:4557877 /UG=Hs.168383 intercellular adhesion molecule 1. (CD54), human rhinovirus receptor /FL=gb:M24283.1 gb:J03132.1 gb:NM_000201.1	NM_000201		NP_000192
202640_s_at	0.029791	gb:NM_003624.1 /DEF=Homo sapiens RAN binding protein 3 (RANBP3), transcript variant RANBP3-a, mRNA. /FEA=mRNA /GEN=RANBP3 /PROD=RAN binding protein 3, isoform RANBP3-a /DB_XREF=gi:4506408 /UG=Hs.176657 RAN binding protein 3 /FL=gb:NM_003624.1	NM_003624		NP_015561

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202641_at	0.002603	gb:NM_004311.1 /DEF=Homo sapiens ADP-ribosylation factor-like 3 (ARL3), mRNA. /FEA=mRNA /GEN=ARL3 /PROD=ADP-ribosylation factor-like 3 /DB_XREF=gi:4757773 /UG=Hs.182215 ADP-ribosylation factor-like 3 /FL=gb:U07151.1 gb:NM_004311.1	NM_004311		NP_004302
202642_s_at	0.036689	gb:NM_003496.1 /DEF=Homo sapiens transformation transcription domain-associated protein (TRRAP), mRNA. /FEA=mRNA /GEN=TRRAP /PROD=transformation transcription domain-associated protein /DB_XREF=gi:4507690 /UG=Hs.203952 transformation transcription domain-associated protein /FL=gb:AF076974.1 gb:NM_003496.1	NM_003496		NP_003487
202645_s_at	0.007652	gb:NM_000244.1 /DEF=Homo sapiens multiple endocrine neoplasia I (MEN1), mRNA. /FEA=mRNA /GEN=MEN1 /PROD=multiple endocrine neoplasia I /DB_XREF=gi:4557744 /UG=Hs.24297 multiple endocrine neoplasia I /FL=gb:U93236.1 gb:NM_000244.1	NM_000244		NP_570716
202649_x_at	0.002603	gb:NM_001022.1 /DEF=Homo sapiens ribosomal protein S19 (RPS19), mRNA. /FEA=mRNA /GEN=RPS19 /PROD=ribosomal protein S19 /DB_XREF=gi:4506694 /UG=Hs.298262 ribosomal protein S19 /FL=gb:BC000023.1 gb:M81757.1 gb:NM_001022.1	NM_001022		NP_001013

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202650_s_at	0.006043	gb:NM_014738.1 /DEF=Homo sapiens KIAA0195 gene product (KIAA0195), mRNA. /FEA=mRNA /GEN=KIAA0195 /PROD=KIAA0195 gene product /DB_XREF=gi:7661985 /UG=Hs.301132 KIAA0195 gene product /FL=gb:D83779.1 gb:NM_014738.1	NM_014738		NP_055553
202651_at	7.25E-04	gb:NM_014873.1 /DEF=Homo sapiens KIAA0205 gene product (KIAA0205), mRNA. /FEA=mRNA /GEN=KIAA0205 /PROD=KIAA0205 gene product /DB_XREF=gi:7661995 /UG=Hs.3610 KIAA0205 gene product /FL=gb:D86960.1 gb:NM_014873.1	NM_014873		NP_055688
202653_s_at	0.03947	Consensus includes gb:BC003404.1 /DEF=Homo sapiens, hypothetical protein DKFZp586F1122 similar to axotrophin, clone IMAGE:3449089, mRNA, partial cds. /FEA=mRNA /PROD=hypothetical protein DKFZp586F1122 similar to axotrophin /DB_XREF=gi:13097302 /UG=Hs.5306 hypothetical protein DKFZp586F1122 similar to axotrophin /FL=gb:NM_022826.1	BC003404		NP_073737

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
202654_x_at	0.003965	gb:NM_022826.1 /DEF=Homo sapiens hypothetical protein DKFZp586F1122 similar to axotrophin (DKFZP586F1122), mRNA. /FEA=mRNA /GEN=DKFZP586F1122 /PROD=hypothetical protein DKFZp586F1122 similar to axotrophin /DB_XREF=gi:12383065 /UG=Hs.5306 hypothetical protein DKFZp586F1122 similar to axotrophin /FL=gb:NM_022826.1	NM_022826		NP_073737
202656_s_at	0.003842	transcriptional regulator interacting with the PHS-bromodomain 2	BG107456	Hs.77293	NP_055570
202657_s_at	7.63E-04	gb:NM_014755.1 /DEF=Homo sapiens KIAA0127 gene product (KIAA0127), mRNA. /FEA=mRNA /GEN=KIAA0127 /PROD=KIAA0127 gene product /DB_XREF=gi:7661925 /UG=Hs.77293 KIAA0127 gene product /FL=gb:D50917.1 gb:NM_014755.1	NM_014755		NP_055570
202658_at	0.044764	gb:NM_003846.1 /DEF=Homo sapiens peroxisomal biogenesis factor 11B (PEX11B), mRNA. /FEA=mRNA /GEN=PEX11B /PROD=peroxisomal biogenesis factor 11B /DB_XREF=gi:4505718 /UG=Hs.83023 peroxisomal biogenesis factor 11B /FL=gb:AF093670.1 gb:AB018080.1 gb:NM_003846.1	NM_003846		NP_003837
202660_at	0.005574	inositol 1,4,5-triphosphate receptor, type 2	AA834576	Hs.238272	

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202662_s_at	0.024403	gb:NM_002223.1 /DEF=Homo sapiens inositol 1,4,5-triphosphate receptor, type 2 (ITPR2), mRNA. /FEA=mRNA /GEN=ITPR2 /PROD=inositol 1,4,5-triphosphate receptor, type 2 /DB_XREF=gi:4504792 /UG=Hs.238272 inositol 1,4,5-triphosphate receptor, type 2 /FL=gb:D26350.1 gb:NM_002223.1	NM_002223		
202663_at	0.004987	Wiskott-Aldrich syndrome protein interacting protein	AI005043	Hs.379203	
202665_s_at	0.024403	gb:NM_003387.2 /DEF=Homo sapiens Wiskott-Aldrich syndrome protein interacting protein (WASPIP), mRNA. /FEA=mRNA /GEN=WASPIP /PROD=WASP-interacting protein /DB_XREF=gi:8400739 /UG=Hs.24143 Wiskott-Aldrich syndrome protein interacting protein /FL=gb:NM_003387.2	NM_003387		NP_003378
202670_at	7.63E-04	mitogen-activated protein kinase kinase 1	AI571419	Hs.3446	NP_002746
202671_s_at	0.001479	gb:NM_003681.1 /DEF=Homo sapiens pyridoxal (pyridoxine, vitamin B6) kinase (PDXK), mRNA. /FEA=mRNA /GEN=PDXK /PROD=pyridoxal kinase /DB_XREF=gi:4505700 /UG=Hs.38041 pyridoxal (pyridoxine, vitamin B6) kinase /FL=gb:BC000123.1 gb:U89606.1 gb:NM_003681.1	NM_003681		NP_003672

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202675_at	7.63E-04	gb:NM_003000.1 /DEF=Homo sapiens succinate dehydrogenase complex, subunit B, iron sulfur (lp) (SDHB), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=SDHB /PROD=succinate dehydrogenase complex, subunit B, iron sulfur (lp) /DB_XREF=gi:9257241 /UG=Hs.64 succinate dehydrogenase complex, subunit B, iron sulfur (lp) /FL=gb:U17248.1 gb:NM_003000.1	NM_003000		NP_002991
202678_at	7.63E-04	gb:NM_004492.1 /DEF=Homo sapiens general transcription factor IIA, 2 (12kD subunit) (GTF2A2), mRNA. /FEA=mRNA /GEN=GTF2A2 /PROD=general transcription factor IIA, 2 (12kD subunit) /DB_XREF=gi:4758485 /UG=Hs.76362 general transcription factor IIA, 2 (12kD subunit) /FL=gb:BC000287.1 gb:BC001919.1 gb:U21242.1 gb:NM_004492.1 gb:U14193.1	NM_004492		NP_004483
202680_at	7.83E-04	gb:NM_002095.1 /DEF=Homo sapiens general transcription factor IIE, polypeptide 2 (beta subunit, 34kD) (GTF2E2), mRNA. /FEA=mRNA /GEN=GTF2E2 /PROD=general transcription factor IIE, polypeptide 2(beta subunit, 34kD) /DB_XREF=gi:4504194 /UG=Hs.77100 general transcription factor IIE, polypeptide 2 (beta subunit, 34kD) /FL=gb:NM_002095.1	NM_002095		NP_002086

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
202687_s_at	0.003976	gb:U57059.1 /DEF=Homo sapiens Apo-2 ligand mRNA, complete cds. /FEA=mRNA /PROD=Apo-2 ligand /DB_XREF=gi:1336207 /UG=Hs.83429 tumor necrosis factor (ligand) superfamily, member 10 /FL=gb:U37518.1 gb:U57059.1 gb:NM_003810.1	U57059		NP_003801
202688_at	0.022601	gb:NM_003810.1 /DEF=Homo sapiens tumor necrosis factor (ligand) superfamily, member 10 (TNFSF10), mRNA. /FEA=mRNA /GEN=TNFSF10 /PROD=tumor necrosis factor (ligand) superfamily, member 10 /DB_XREF=gi:4507592 /UG=Hs.83429 tumor necrosis factor (ligand) superfamily, member 10 /FL=gb:U37518.1 gb:U57059.1 gb:NM_003810.1	NM_003810		NP_003801
202690_s_at	0.020369	gb:BC001721.1 /DEF=Homo sapiens, small nuclear ribonucleoprotein D1 polypeptide (16kD), clone MGC:2222, mRNA, complete cds. /FEA=mRNA /PROD=small nuclear ribonucleoprotein D1 polypeptide(16kD) /DB_XREF=gi:12804598 /UG=Hs.86948 small nuclear ribonucleoprotein D1 polypeptide (16kD) /FL=gb:BC001721.1 gb:J03798.1 gb:NM_006938.1	BC001721		NP_008869

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	D scription	Gene Accession No.	Unigen Accession No.	Protein Accession No.
202696_at	7.63E-04	gb:NM_005109.1 /DEF=Homo sapiens oxidative-stress responsive 1 (OSR1), mRNA. /FEA=mRNA /GEN=OSR1 /PROD=oxidative-stress responsive 1 /DB_XREF=gi:4826877 /UG=Hs.95220 oxidative-stress responsive 1 /FL=gb:AB017642.1 gb:NM_005109.1 gb:AB029024.1	NM_005109		NP_005100
202697_at	7.76E-04	gb:NM_007006.1 /DEF=Homo sapiens cleavage and polyadenylation specific factor 5, 25 kD subunit (CPSF5), mRNA. /FEA=mRNA /GEN=CPSF5 /PROD=cleavage and polyadenylation specific factor 5,25 kD subunit /DB_XREF=gi:5901925 /UG=Hs.9605 cleavage and polyadenylation specific factor 5, 25 kD subunit /FL=gb:BC001403.1 gb:NM_007006.1	NM_007006		NP_008937
202700_s_at	0.019717	gb:NM_014698.1 /DEF=Homo sapiens KIAA0792 gene product (KIAA0792), mRNA. /FEA=mRNA /GEN=KIAA0792 /PROD=KIAA0792 gene product /DB_XREF=gi:7662307 /UG=Hs.119387 KIAA0792 gene product /FL=gb:AB018335.1 gb:NM_014698.1	NM_014698		NP_055513

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202708_s_at	0.018901	gb:NM_003528.1 /DEF=Homo sapiens H2B histone family, member Q (H2BFQ), mRNA. /FEA=mRNA /GEN=H2BFQ /PROD=H2B histone family, member Q /DB_XREF=gi:4504276 /UG=Hs.2178 H2B histone family, member Q /FL=gb:NM_003528.1	NM_003528		NP_003519
202719_s_at	0.040906	gb:BC001451.1 /DEF=Homo sapiens, testin, clone MGC:1146, mRNA, complete cds. /FEA=mRNA /PROD=testin /DB_XREF=gi:12655188 /UG=Hs.165986 testin /FL=gb:AF245356.1 gb:AF245357.1 gb:BC001451.1 gb:NM_015641.1	BC001451		NP_690042
202720_at	0.037453	gb:NM_015641.1 /DEF=Homo sapiens testin (DKFZP586B2022), mRNA. /FEA=mRNA /GEN=DKFZP586B2022 /PROD=testin. /DB_XREF=gi:7661665 /UG=Hs.165986 testin /FL=gb:AF245356.1 gb:AF245357.1 gb:BC001451.1 gb:NM_015641.1	NM_015641		NP_690042
202725_at	0.001179	gb:NM_000937.1 /DEF=Homo sapiens polymerase (RNA) II (DNA directed) polypeptide A (220kD) (POLR2A), mRNA. /FEA=mRNA /GEN=POLR2A /PROD=polymerase (RNA) II (DNA directed) polypeptide A(220kD) /DB_XREF=gi:4505938 /UG=Hs.171880 polymerase (RNA) II (DNA directed) polypeptide A (220kD) /FL=gb:NM_000937.1	NM_000937		NP_000928

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202727_s_at	0.001719	gb:NM_000416.1 /DEF=Homo sapiens interferon gamma receptor 1 (IFNGR1), mRNA. /FEA=mRNA /GEN=IFNGR1 /PROD=interferon gamma receptor 1 /DB_XREF=gi:4557879 /UG=Hs.180866 interferon gamma receptor 1 /FL=gb:BC005333.1 gb:J03143.1 gb:NM_000416.1	NM_000416		NP_000407
202731_at	0.01747	gb:NM_014456.1 /DEF=Homo sapiens programmed cell death 4 (PDCD4), mRNA. /FEA=mRNA /GEN=PDCD4 /PROD=programmed cell death 4 /DB_XREF=gi:7657448 /UG=Hs.296251 programmed cell death 4 /FL=gb:U96628.1 gb:NM_014456.1	NM_014456		NP_663314
202738_s_at	0.018248	phosphorylase kinase, beta	BG149218	Hs.78060	NP_000284
202742_s_at	0.00643	gb:NM_002731.1 /DEF=Homo sapiens protein kinase, cAMP-dependent, catalytic, beta (PRKACB), mRNA. /FEA=mRNA /GEN=PRKACB /PROD=protein kinase, cAMP-dependent, catalytic, beta /DB_XREF=gi:4506056 /UG=Hs.87773 protein kinase, cAMP-dependent, catalytic, beta /FL=gb:M34181.1 gb:NM_002731.1	NM_002731		NP_002722
202749_at	0.026836	gb:NM_004627.1 /DEF=Homo sapiens tryptophan rich basic protein (WRB), mRNA. /FEA=mRNA /GEN=WRB /PROD=tryptophan rich basic protein /DB_XREF=gi:4759325 /UG=Hs.198308 tryptophan rich basic protein /FL=gb:NM_004627.1	NM_004627		NP_004618

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202750_s_at	0.013551	gb:AL080147.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434B194 (from clone DKFZp434B194); complete cds. /FEA=mRNA /GEN=DKFZp434B194 /PROD=hypothetical protein /DB_XREF=gi:5262598 /UG=Hs.20225 tuftelin-interacting protein. /FL=gb:AL080147.1 gb:NM_012143.1	AL080147		NP_036275
202753_at	0.018534	gb:NM_014814.1 /DEF=Homo sapiens KIAA0107 gene product (KIAA0107), mRNA. /FEA=mRNA /GEN=KIAA0107 /PROD=KIAA0107 gene product /DB_XREF=gi:7661913 /UG=Hs.23488 KIAA0107 gene product /FL=gb:BC000630.1 gb:BC000904.2 gb:D14663.1 gb:AF215935.1 gb:NM_014814.1	NM_014814		NP_055629
202758_s_at	0.017898	gb:NM_003721.1 /DEF=Homo sapiens regulatory factor X-associated ankyrin-containing protein (RFXANK), mRNA. /FEA=mRNA /GEN=RFXANK /PROD=regulatory factor X-associated ankyrin-containing protein /DB_XREF=gi:4506498 /UG=Hs.296776 regulatory factor X-associated ankyrin-containing protein /FL=gb:AF094760.1 gb:AF105427.1 gb:NM_003721.1 gb:AF077196.1	NM_003721		NP_604389

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202761_s_at	0.043116	gb:NM_015180.1 /DEF=Homo sapiens synaptic nuclei expressed gene 2; KIAA1011 protein (KIAA1011), mRNA. /FEA=mRNA /GEN=KIAA1011 /PROD=KIAA1011 protein /DB_XREF=gi:11056019 /UG=Hs.57749 synaptic nuclei expressed gene 2; KIAA1011 protein /FL=gb:NM_015180.1 gb:AL080133.1	NM_015180		
202764_at	0.004814	gb:NM_003156.1 /DEF=Homo sapiens stromal interaction molecule 1 (STIM1), mRNA. /FEA=mRNA /GEN=STIM1 /PROD=stromal interaction molecule 1 /DB_XREF=gi:4507268 /UG=Hs.74597 stromal interaction molecule 1 /FL=gb:U52426.1 gb:NM_003156.1	NM_003156		NP_003147
202767_at	0.003009	gb:NM_001610.1 /DEF=Homo sapiens acid phosphatase 2, lysosomal (ACP2), mRNA. /FEA=mRNA /GEN=ACP2 /PROD=lysosomal acid phosphatase 2 precursor /DB_XREF=gi:4557009 /UG=Hs.75589 acid phosphatase 2, lysosomal /FL=gb:BC003160.1 gb:NM_001610.1	NM_001610		NP_001601
202770_s_at	0.002692	gb:NM_004354.1 /DEF=Homo sapiens cyclin G2 (CCNG2), mRNA. /FEA=mRNA /GEN=CCNG2 /PROD=cyclin G2 /DB_XREF=gi:4757935 /UG=Hs.79069 cyclin G2 /FL=gb:U47414.1 gb:NM_004354.1	NM_004354		NP_004345
202774_s_at	0.047091	splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila)	AI023864	Hs.84229	NP_689421

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202777_at	7.25E-04	gb:NM_007373.1 /DEF=Homo sapiens suppressor of clear, C. elegans, homolog of (SHOC2), mRNA. /FEA=mRNA /GEN=SHOC2 /PROD=suppressor of clear, C. elegans, homolog of /DB_XREF=gi:6677944 /UG=Hs.104315 soc-2 (suppressor of clear, C.elegans) homolog /FL=gb:AF068920.1 gb:AF054828.1 gb:AB020669.1 gb:NM_007373.1	NM_007373		NP_031399
202782_s_at	0.025377	gb:NM_016532.1 /DEF=Homo sapiens SKIP for skeletal muscle and kidney enriched inositol phosphatase (LOC51763), mRNA. /FEA=mRNA /GEN=LOC51763 /PROD=SKIP for skeletal muscle and kidney enriched inositol phosphatase /DB_XREF=gi:7706564 /UG=Hs.178347 SKIP for skeletal muscle and kidney enriched inositol phosphatase /FL=gb:AB036829.1 gb:NM_016532.1	NM_016532		NP_570122
202788_at	0.00109	gb:NM_004635.1 /DEF=Homo sapiens mitogen-activated protein kinase-activated protein kinase 3 (MAPKAPK3), mRNA. /FEA=mRNA /GEN=MAPKAPK3 /PROD=mitogen-activated protein kinase-activated protein kinase 3 /DB_XREF=gi:4758699 /UG=Hs.227789 mitogen-activated protein kinase-activated protein kinase 3 /FL=gb:U09578.1 gb:U43784.1 gb:BC001662.1 gb:NM_004635.1	NM_004635		NP_004626

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
202794_at	0.006864	gb:NM_002194.2 /DEF=Homo sapiens inositol polyphosphate-1-phosphatase (INPP1), mRNA. /FEA=mRNA /GEN=INPP1 /PROD=inositol polyphosphate-1-phosphatase /DB_XREF=gi:4755138 /UG=Hs.32309 inositol polyphosphate-1-phosphatase. /FL=gb:L08488.1 gb:NM_002194.2	NM_002194		NP_002185
202798_at	0.024403	gb:NM_006323.1 /DEF=Homo sapiens SEC24 (S. cerevisiae) related gene family, member B (SEC24B), mRNA. /FEA=mRNA /GEN=SEC24B /PROD=SEC24 (S. cerevisiae) related gene family, member B /DB_XREF=gi:5454045 /UG=Hs.7239 SEC24 (S. cerevisiae) related gene family, member B /FL=gb:NM_006323.1	NM_006323		NP_006314
2028_s_at	0.012808	E2F transcription factor 1	NM_005225	Hs.96055	NP_005216
202807_s_at	0.004814	gb:NM_005488.1 /DEF=Homo sapiens target of myb1 (chicken) homolog (TOM1), mRNA. /FEA=mRNA /GEN=TOM1 /PROD=target of myb1 (chicken) homolog /DB_XREF=gi:4885636 /UG=Hs.9482 target of myb1 (chicken) homolog /FL=gb:NM_005488.1	NM_005488		NP_005479

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202810_at	0.049173	gb:NM_004147.1 /DEF=Homo sapiens developmentally regulated GTP-binding protein 1 (DRG1), mRNA. /FEA=mRNA /GEN=DRG1 /PROD=developmentally regulated GTP-binding protein 1 /DB_XREF=gi:4758795 /UG=Hs.115242 developmentally regulated GTP-binding protein 1 /FL=gb:AF078103.1 gb:NM_004147.1	NM_004147		NP_004138
202811_at	0.002749	gb:NM_006463.1 /DEF=Homo sapiens associated molecule with the SH3 domain of STAM (AMSH), mRNA. /FEA=mRNA /GEN=AMSH /PROD=associated molecule with the SH3 domain of STAM /DB_XREF=gi:5453544 /UG=Hs.12479 associated molecule with the SH3 domain of STAM /FL=gb:U73522.1 gb:NM_006463.1	NM_006463		NP_006454
202814_s_at	0.044966	HMBA-inducible	AW193511	Hs.15299	NP_006451
202824_s_at	0.001852	gb:NM_005648.1 /DEF=Homo sapiens transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C) (TCEB1), mRNA. /FEA=mRNA /GEN=TCEB1 /PROD=elongin C /DB_XREF=gi:5032160 /UG=Hs.184693 transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C) /FL=gb:NM_005648.1 gb:L34587.1	NM_005648		NP_005639
202829_s_at	7.76E-04	gb:NM_005638.1 /DEF=Homo sapiens synaptobrevin-like 1 (SYBL1), mRNA. /FEA=mRNA /GEN=SYBL1 /PROD=synaptobrevin-like 1 /DB_XREF=gi:5032136 /UG=Hs.24167 synaptobrevin-like 1 /FL=gb:NM_005638.1	NM_005638		NP_005629

Genes Corresponding To Differentially Expressed Genes in Figur 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202833_s_at	7.25E-04	gb:NM_000295.1 /DEF=Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1 (SERPINA1), mRNA. /FEA=mRNA /GEN=SERPINA1 /PROD=serine (or cysteine) proteinase inhibitor, cladeA (alpha-1 antiproteinase, antitrypsin), member 1 /DB_XREF=gi:4505792 /UG=Hs.297681 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1. /FL=gb:AF130068.1 gb:M11465.1 gb:K01396.1 gb:NM_000295.1	NM_000295		NP_000286
202836_s_at	0.003965	gb:NM_006701.1 /DEF=Homo sapiens similar to S. pombe dim1+ (DIM1), mRNA. /FEA=mRNA /GEN=DIM1 /PROD=similar to S. pombe dim1+ /DB_XREF=gi:5729801 /UG=Hs.5074 similar to S. pombe dim1+ /FL=gb:BC001046.1 gb:AF023611.1 gb:NM_006701.1 gb:AF146373.1	NM_006701		NP_006692
202845_s_at	0.001719	gb:NM_006788.1 /DEF=Homo sapiens ralA binding protein 1 (RALBP1), mRNA. /FEA=mRNA /GEN=RALBP1 /PROD=ralA binding protein 1 /DB_XREF=gi:5803144 /UG=Hs.75447 ralA binding protein 1 /FL=gb:NM_006788.1 gb:L42542.1	NM_006788		NP_006779
202848_s_at	0.03359	G protein-coupled receptor kinase 6	BG423052	Hs.76297	NP_002073

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202852_s_at	0.002786	gb:NM_024666.1 /DEF=Homo sapiens hypothetical protein FLJ11506 (FLJ11506), mRNA. /FEA=mRNA /GEN=FLJ11506 /PROD=hypothetical protein FLJ11506 /DB_XREF=gi:13375923 /UG=Hs.77703 hypothetical protein FLJ11506 /FL=gb:AL136715.1 gb:NM_024666.1	NM_024666		NP_078942
202854_at	7.63E-04	gb:NM_000194.1 /DEF=Homo sapiens hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan syndrome) (HPRT1), mRNA. /FEA=mRNA /GEN=HPRT1 /PROD=hypoxanthine phosphoribosyltransferase 1 /DB_XREF=gi:4504482 /UG=Hs.82314 hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan syndrome) /FL=gb:BC000578.1 gb:M31642.1 gb:NM_000194.1	NM_000194		NP_000185
202859_x_at	0.009678	gb:NM_000584.1 /DEF=Homo sapiens interleukin 8 (IL8), mRNA. /FEA=mRNA /GEN=IL8 /PROD=interleukin 8 /DB_XREF=gi:10834977 /UG=Hs.624 interleukin 8 /FL=gb:NM_000584.1 gb:M17017.1 gb:M26383.1	NM_000584		NP_000575
202872_at	9.85E-04	Consensus includes gb:AW024925 /FEA=EST /DB_XREF=gi:5878455 /DB_XREF=est:wu92g09.x1 /CLONE=IMAGE:2527552 /UG=Hs.86905 ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 42kD /FL=gb:NM_001695.1	NM_001695		NP_001686

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202873_at	0.03002	Consensus includes gb:BF034973 /FEA=EST /DB_XREF=gi:10742685 /DB_XREF=est:601459039F1 /CLONE=IMAGE:3862832 /UG=Hs.86905 ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD /FL=gb:NM_001695.1	NM_001695		NP_001686
202874_s_at	0.007591	gb:NM_001695.1 /DEF=Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD (ATP6C), mRNA. /FEA=mRNA /GEN=ATP6C /PROD=ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD /DB_XREF=gi:4502314 /UG=Hs.86905 ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD /FL=gb:NM_001695.1	NM_001695		NP_001686
202878_s_at	0.00121	gb:NM_012072.2 /DEF=Homo sapiens complement component C1q receptor (C1QR), mRNA. /FEA=mRNA /GEN=C1QR /PROD=complement component C1q receptor /DB_XREF=gi:11496985 /UG=Hs.97199 complement component C1q receptor /FL=gb:NM_012072.2 gb:U94333.1	NM_012072		NP_036204

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202880_s_at	0.009329	gb:NM_004762.1 /DEF=Homo sapiens pleckstrin homology, Sec7 and coiledcoil domains 1(cytohesin 1) (PSCD1), transcript variant 1; mRNA. /FEA=mRNA /GEN=PSCD1 /PROD=cytohesin 1, isoform 1 /DB_XREF=gi:4758963 /UG=Hs.1050 pleckstrin homology, Sec7 and coiledcoil domains 1(cytohesin 1) /FL=gb:M85169.1 gb:NM_004762.1 gb:NM_017456.1	NM_004762		NP_059430
202882_x_at	0.022439	gb:NM_016167.2 /DEF=Homo sapiens retinoic acid repressible protein (RAR-1); mRNA. /FEA=mRNA /GEN=RARG-1 /PROD=retinoic acid repressible protein /DB_XREF=gi:13569842 /UG=Hs.106346 retinoic acid repressible protein /FL=gb:AF172066.2 gb:NM_016167.2	NM_016167		NP_057251
202888_s_at	0.031624	gb:NM_001150.1 /DEF=Homo sapiens alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150) (ANPEP), mRNA. /FEA=mRNA /GEN=ANPEP /PROD=membrane alanine aminopeptidase precursor /DB_XREF=gi:4502094 /UG=Hs.1239 alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150) /FL=gb:M22324.1 gb:NM_001150.1	NM_001150		NP_001141

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202891_at	7.63E-04	gb:NM_005600.1 /DEF=Homo sapiens nitrilase 1 (NIT1), mRNA: /FEA=mRNA /GEN=NIT1 /PROD=nitrilase 1 /DB_XREF=gi:5031946 /UG=Hs.146406 nitrilase 1 /FL=gb:AF069987.1 gb:NM_005600.1	NM_005600		NP_005591
202897_at	7.25E-04	gb:AB023430.1 /DEF=Homo sapiens Bit mRNA, complete cds. /FEA=mRNA /GEN=Bit /DB_XREF=gi:6518912 /UG=Hs.156114 protein tyrosine phosphatase, non-receptor type substrate 1 /FL=gb:D86043.1 gb:NM_004648.1 gb:AB023430.1	AB023430		NP_542970
202907_s_at	0.005347	gb:NM_002485.2 /DEF=Homo sapiens Nijmegen breakage syndrome 1 (nibrin) (NBS1), mRNA: /FEA=mRNA /GEN=NBS1 /PROD=nibrin /DB_XREF=gi:6996019 /UG=Hs.25812 Nijmegen breakage syndrome 1 (nibrin) /FL=gb:AF058696.1 gb:AF051334.1 gb:NM_002485.2	NM_002485		NP_002476
202918_s_at	0.024696	gb:AF151853.1 /DEF=Homo sapiens CGI-95 protein mRNA, complete cds. /FEA=mRNA /PROD=CGI-95 protein /DB_XREF=gi:4929658 /UG=Hs.107942 DKFZP564M112 protein /FL=gb:AB015441.1 gb:BC005237.1 gb:AF151853.1 gb:AL080070.1 gb:NM_015387.1	AF151853		NP_056202

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202925_s_at	0.003373	gb:NM_002657.2 /DEF=Homo sapiens pleiomorphic adenoma gene-like 2 (PLAGL2), mRNA. /FEA=mRNA /GEN=PLAGL2 /PROD=pleiomorphic adenoma gene-like 2 /DB_XREF=gi:6031195 /UG=Hs.154104 pleiomorphic adenoma gene-like 2 /FL=gb:AF006005.1 gb:NM_002657.2	NM_002657		NP_002648
202934_at	7.63E-04	hexokinase 2	A1761561	Hs.198427	NP_000180
202939_at	0.001406	gb:NM_005857.1 /DEF=Homo sapiens zinc metalloproteinase, STE24 (yeast, homolog) (ZMPSTE24), mRNA. /FEA=mRNA /GEN=ZMPSTE24 /PROD=zinc metalloproteinase, STE24 (yeast, homolog) /DB_XREF=gi:5032128 /UG=Hs.25846 zinc metalloproteinase, STE24 (yeast, homolog) /FL=gb:AB016068.1 gb:AF064867.1 gb:NM_005857.1	NM_005857		NP_005848
202941_at	0.003976	gb:NM_021074.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) flavoprotein 2 (24kD) (NDUFV2), mRNA. /FEA=mRNA /GEN=NDUFV2 /PROD=NADH dehydrogenase (ubiquinone) flavoprotein 2(24kD) /DB_XREF=gi:10835024 /UG=Hs.51299 NADH dehydrogenase (ubiquinone) flavoprotein 2 (24kD) /FL=gb:NM_021074.1 gb:BC001632.1 gb:M22538.1	NM_021074		NP_066552

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
202944_at	0.025675	gb:NM_000262.1 /DEF=Homo sapiens N-acetylgalactosaminidase, alpha- (NAGA), mRNA. /FEA=mRNA /GEN=NAGA /PROD=alpha-N-acetylgalactosaminidase precursor /DB_XREF=gi:4557780 /UG=Hs.75372 N-acetylgalactosaminidase, alpha- /FL=gb:BC000095.1 gb:M62783.1 gb:M38083.1 gb:NM_000262.1	NM_000262		NP_000253
202946_s_at	0.001231	gb:NM_014962.1 /DEF=Homo sapiens KIAA0952 protein (KIAA0952), mRNA. /FEA=mRNA /GEN=KIAA0952 /PROD=KIAA0952 protein /DB_XREF=gi:7662401 /UG=Hs.7935 KIAA0952 protein /FL=gb:AB023169.1 gb:NM_014962.1	NM_014962		NP_852108
202947_s_at	0.001433	gb:NM_002101.2 /DEF=Homo sapiens glycophorin C (Gerbich blood group) (GYPC), transcript variant 1, mRNA. /FEA=mRNA /GEN=GYPC /PROD=glycophorin C, isoform 1 /DB_XREF=gi:8051606 /UG=Hs.81994 glycophorin C (Gerbich blood group) /FL=gb:M36284.1 gb:M11802.1 gb:NM_002101.2	NM_002101		NP_058131

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202948_at	7.85E-04	gb:NM_000877.1 /DEF=Homo sapiens interleukin 1 receptor, type I (IL1R1), mRNA. /FEA=mRNA /GEN=IL1R1 /PROD=interleukin 1 receptor, type I /DB_XREF=gi:4504658 /UG=Hs.82112 interleukin 1 receptor, type I /FL=gb:M27492.1 gb:NM_000877.1	NM_000877		NP_000868
202954_at	0.023856	gb:NM_007019.1 /DEF=Homo sapiens ubiquitin carrier protein E2-C (UBCH10), mRNA. /FEA=mRNA /GEN=UBCH10 /PROD=ubiquitin carrier protein E2-C /DB_XREF=gi:5902145 /UG=Hs.93002 ubiquitin carrier protein E2-C /FL=gb:U73379.1 gb:NM_007019.1	NM_007019		NP_008950
202956_at	0.002109	gb:NM_006421.2 /DEF=Homo sapiens brefeldin A-inhibited guanine nucleotide-exchange protein 1 (BIG1), mRNA. /FEA=mRNA /GEN=BIG1 /PROD=brefeldin A-inhibited guanine nucleotide-exchange protein 1 /DB_XREF=gi:6715588 /UG=Hs.94631 brefeldin A-inhibited guanine nucleotide-exchange protein 1 /FL=gb:AF084520.1 gb:AF111162.1 gb:NM_006421.2	NM_006421		NP_006412

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unig n Accession No.	Protein Acc ssion No.
202960_s_at	0.030968	gb:NM_000255.1 /DEF=Homo sapiens methylmalonyl Coenzyme A mutase (MUT), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=MUT /PROD=methylmalonyl Coenzyme A mutase precursor /DB_XREF=gi:4557766 /UG=Hs.155212 methylmalonyl Coenzyme A mutase /FL=gb:M65131.1 gb:NM_000255.1	NM_000255		NP_000246
202961_s_at	0.003965	gb:NM_004889.1 /DEF=Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, subunit f, isoform 2 (ATP5J2), mRNA. /FEA=mRNA /GEN=ATP5J2 /PROD=ATP synthase, H+ transporting, mitochondrial F0 complex, subunit f, isoform 2 /DB_XREF=gi:4757811 /UG=Hs.155751 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit f, isoform 2 /FL=gb:BC003678.1 gb:AF047436.1 gb:NM_004889.1	NM_004889		NP_004880
202968_s_at	0.028603	Consensus includes gb:Y09216.1 /DEF=H.sapiens mRNA for protein kinase, Dyrk2. /FEA=mRNA /GEN=Dyrk2 /DB_XREF=gi:1666065 /UG=Hs.173135 dual-specificity tyrosine-(Y)-phosphorylation-regulated kinase 2 /FL=gb:NM_006482.1	Y09216		NP_006473

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Prot in Accession No.
202969_at	0.005652	Consensus includes gb:AI216690 /FEA=EST /DB_XREF=gi:3785731 /DB_XREF=est:qg66h07.x1 /CLONE=IMAGE:1840189 /UG=Hs.173135 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 /FL=gb:NM_006482.1	Y09216		NP_006473
202971_s_at	0.018693	gb:NM_006482.1 /DEF=Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 (DYRK2), transcript variant 2, mRNA. /FEA=mRNA /GEN=DYRK2 /PROD=dual-specificity tyrosine- (Y)-phosphorylationregulated kinase 2 isoform 2 /DB_XREF=gi:5922003 /UG=Hs.173135 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 /FL=gb:NM_006482.1	NM_006482		NP_006473
202974_at	6.66E-04	gb:NM_002436.2 /DEF=Homo sapiens membrane protein, palmitoylated 1 (55kD) (MPP1), mRNA. /FEA=mRNA /GEN=MPP1 /PROD=palmitoylated membrane protein 1 /DB_XREF=gi:6006024 /UG=Hs.1861 membrane protein, palmitoylated 1 (55kD) /FL=gb:BC002392.1 gb:M64925.1 gb:NM_002436.2	NM_002436		NP_002427

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202990_at	7.25E-04	gb:NM_002863.1 /DEF=Homo sapiens phosphorylase, glycogen; liver (Hers disease, glycogen storage disease type VI) (PYGL), mRNA. /FEA=mRNA /GEN=PYGL /PROD=phosphorylase, glycogen; liver (Hers disease, glycogen storage disease type VI) /DB_XREF=gi:4506352 /UG=Hs.771 phosphorylase, glycogen; liver (Hers disease, glycogen storage disease type VI) /FL=gb:M14636.1 gb:AF066858.1 gb:AF046785.1 gb:NM_002863.1	NM_002863		NP_002854
202995_s_at	0.003288	gb:NM_006486.1 /DEF=Homo sapiens fibulin 1 (FBLN1), transcript variant D, mRNA. /FEA=mRNA /GEN=FBLN1 /PROD=fibulin 1 isoform D /DB_XREF=gi:5922008 /UG=Hs.79732 fibulin 1 /FL=gb:U01244.1 gb:NM_006486.1	NM_006486		NP_006478
203005_at	7.25E-04	gb:NM_002342.1 /DEF=Homo sapiens lymphotoxin beta receptor (TNFR superfamily, member 3 (LTBR), mRNA. /FEA=mRNA /GEN=LTBR /PROD=lymphotoxin B receptor /DB_XREF=gi:4505038 /UG=Hs.1116 lymphotoxin beta receptor (TNFR superfamily, member 3 /FL=gb:L04270.1 gb:NM_002342.1	NM_002342		NP_002333

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
203008_x_at	0.02217	gb:NM_005783.1 /DEF=Homo sapiens ATP binding protein associated with cell differentiation (APACD), mRNA. /FEA=mRNA /GEN=APACD /PROD=ATP binding protein associated with celldifferentiation /DB_XREF=gi:5031582 /UG=Hs.153884 ATP binding protein associated with cell differentiation /FL=gb:AB006679.1 gb:NM_005783.1	NM_005783		NP_005774
203011_at	0.005574	gb:NM_005536.2 /DEF=Homo sapiens inositol(myo)-1(or 4)-monophosphatase 1 (IMPA1), mRNA. /FEA=mRNA /GEN=IMPA1 /PROD=inositol(myo)-1(or 4)-monophosphatase 1 /DB_XREF=gi:8393607 /UG=Hs.171776 inositol(myo)-1(or 4)-monophosphatase 1 /FL=gb:AF042729.2 gb:NM_005536.2	NM_005536		NP_005527
203021_at	0.00121	gb:NM_003064.1 /DEF=Homo sapiens secretory leukocyte protease inhibitor (antileukoproteinase) (SLPI), mRNA. /FEA=mRNA /GEN=SLPI /PROD=secretory leukocyte protease inhibitor(antileukoproteinase) /DB_XREF=gi:4507064 /UG=Hs.251754 secretory leukocyte protease inhibitor (antileukoproteinase) /FL=gb:NM_003066.1 gb:AF114471.1 gb:NM_003064.1	NM_003064		NP_003055

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203025_at	0.022474	gb:NM_003491.1 /DEF=Homo sapiens N-acetyltransferase, homolog of S. cerevisiae ARD1 (ARD1), mRNA. /FEA=mRNA /GEN=ARD1 /PROD=N-acetyltransferase, homolog of S. cerevisiae ARD1 /DB_XREF=gi:10835056 /UG=Hs.333034 N-acetyltransferase, homolog of S. cerevisiae ARD1 /FL=gb:NM_003491.1 gb:BC000308.1	NM_003491		NP_003482
203039_s_at	0.003976	gb:NM_005006.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 1 (75kD) (NADH-coenzyme Q reductase) (NDUFS1), mRNA. /FEA=mRNA /GEN=NDUFS1 /PROD=NADH dehydrogenase (ubiquinone) Fe-S protein 1(75kD) (NADH-coenzyme Q reductase) /DB_XREF=gi:4826855 /UG=Hs.8248 NADH dehydrogenase (ubiquinone) Fe-S protein 1 (75kD). (NADH-coenzyme Q reductase) /FL=gb:NM_005006.1	NM_005006		NP_004997
203041_s_at	7.25E-04	gb:J04183.1 /DEF=Homo sapiens lysosomal membrane glycoprotein-2 (LAMP2), complete cds. /FEA=mRNA /GEN=LAMP2 /PROD=lysosomal membrane glycoprotein-2 /DB_XREF=gi:186929 /UG=Hs.8262 lysosomal-associated membrane protein 2 /FL=gb:J04183.1 gb:NM_002294.1	J04183		NP_054701

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203042_at	0.00109	gb:NM_002294.1 /DEF=Homo sapiens lysosomal-associated membrane protein 2 (LAMP2), transcript variant LAMP2A, mRNA. /FEA=mRNA /GEN=LAMP2 /PROD=lysosomal-associated membrane protein 2precursor /DB_XREF=gi:4504956 /UG=Hs.8262 lysosomal-associated membrane protein 2 /FL=gb:J04183.1 gb:NM_002294.1	NM_002294		NP_054701
203044_at	0.00158	gb:NM_014918.1 /DEF=Homo sapiens KIAA0990 protein (KIAA0990), mRNA. /FEA=mRNA /GEN=KIAA0990 /PROD=KIAA0990 protein /DB_XREF=gi:7662433 /UG=Hs.110488 KIAA0990 protein. /FL=gb:AB023207.1 gb:NM_014918.1	NM_014918		NP_055733
203057_s_at	0.049392	Consensus includes gb:AV724783 /FEA=EST /DB_XREF=gi:10829513 /DB_XREF=est:AV724783 /CLONE=HTBCEG11 /UG=Hs.26719 zinc-finger DNA-binding protein /FL=gb:D45132.1 gb:NM_015866.1	NM_015866		NP_056950
203064_s_at	0.036752	gb:NM_004514.1 /DEF=Homo sapiens interleukin enhancer binding factor 1 (ILF1), mRNA. /FEA=mRNA /GEN=ILF1 /PROD=interleukin enhancer binding factor 1 /DB_XREF=gi:4758599 /UG=Hs.296281 interleukin enhancer binding factor 1 /FL=gb:U58196.1 gb:NM_004514.1	NM_004514		NP_004505

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
203074_at	0.030776	gb:NM_001630.1 /DEF=Homo sapiens annexin A8 (ANXA8), mRNA. /FEA=mRNA /GEN=ANXA8 /PROD=annexin VIII /DB_XREF=gi:4502112 /UG=Hs.87268 annexin A8 /FL=gb:BC004376.1 gb:M81844.1 gb:NM_001630.1	NM_001630		NP_001621
203075_at	0.001835	MAD, mothers against decapentaplegic homolog 2 (Drosophila)	AW151617	Hs.82483	NP_005892
203076_s_at	7.63E-04	gb:U65019.1 /DEF=Human MAD-related protein 2 (MADR2) mRNA; complete cds. /FEA=mRNA /GEN=MADR2 /PROD=MAD-related protein 2 /DB_XREF=gi:1575529 /UG=Hs.82483 MAD (mothers against decapentaplegic, Drosophila) homolog 2 /FL=gb:U59911.1 gb:U68018.1 gb:U65019.1 gb:AF027964.1 gb:NM_005901.1	U65019		NP_005892
203077_s_at	0.027672	gb:NM_005901.1 /DEF=Homo sapiens MAD (mothers against decapentaplegic, Drosophila) homolog 2 (MADH2), mRNA. /FEA=mRNA /GEN=MADH2 /PROD=MAD (mothers against decapentaplegic, Drosophila) homolog 2 /DB_XREF=gi:5174510 /UG=Hs.82483 MAD (mothers against decapentaplegic, Drosophila) homolog 2 /FL=gb:U59911.1 gb:U68018.1 gb:U65019.1 gb:AF027964.1 gb:NM_005901.1	NM_005901		NP_005892

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203082_at	0.031624	gb:NM_014753.1 /DEF=Homo sapiens KIAA0187 gene product (KIAA0187), mRNA. /FEA=mRNA /GEN=KIAA0187 /PROD=KIAA0187 gene product /DB_XREF=gi:7661979 /UG=Hs.10848 KIAA0187 gene product /FL=gb:D80009.1 gb:NM_014753.1	NM_014753		NP_055568
203090_at	0.00158	gb:NM_006923.1 /DEF=Homo sapiens stromal cell-derived factor 2 (SDF2), mRNA. /FEA=mRNA /GEN=SDF2 /PROD=stromal cell-derived factor 2 /DB_XREF=gi:5902073 /UG=Hs.118684 stromal cell-derived factor 2 /FL=gb:BC000500.1 gb:BC001406.1 gb:D50645.1 gb:NM_006923.1	NM_006923		NP_008854
203094_at	0.024932	gb:NM_014628.1 /DEF=Homo sapiens gene predicted from cDNA with a complete coding sequence (KIAA0110), mRNA. /FEA=mRNA /GEN=KIAA0110 /PROD=gene predicted from cDNA with a complete coding sequence /DB_XREF=gi:7661917 /UG=Hs.124 gene predicted from cDNA with a complete coding sequence /FL=gb:BC002904.1 gb:D14811.1 gb:NM_014628.1	NM_014628		NP_055443

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203097_s_at	0.013527	gb:NM_014247.1 /DEF=Homo sapiens PDZ domain containing guanine nucleotide exchange factor(GEF)1 (PDZ-GEF1), mRNA. /FEA=mRNA /GEN=PDZ-GEF1 /PROD=PDZ domain containing guanine nucleotideexchange factor(GEF)1 /DB_XREF=gi:7657260 /UG=Hs.154545 PDZ domain containing guanine nucleotide exchange factor(GEF)1 /FL=gb:AB002311.1 gb:NM_014247.1	NM_014247		NP_055062
203100_s_at	0.034328	gb:NM_004824.1 /DEF=Homo sapiens chromodomain protein, Y chromosome-like (CDYL), mRNA. /FEA=mRNA /GEN=CDYL /PROD=chromodomain protein, Y chromosome-like /DB_XREF=gi:4757969 /UG=Hs.16081 chromodomain protein, Y chromosome-like /FL=gb:AF081258.1 gb:AF081259.1 gb:NM_004824.1	NM_004824		NP_736608
203102_s_at	0.029624	gb:NM_002408.2 /DEF=Homo sapiens mannosyl (alpha-1,6-) glycoprotein beta-1,2-N-acetylglucosaminyltransferase (MGAT2), mRNA. /FEA=mRNA /GEN=MGAT2 /PROD=alpha-1,6-mannosyl-glycoproteinbeta-1,2-N-acetylglucosaminyltransferase /DB_XREF=gi:6031183 /UG=Hs.172195 mannosyl (alpha-1,6-) glycoprotein beta-1,2-N-acetylglucosaminyltransferase /FL=gb:NM_002408.2	NM_002408		NP_002399

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203107_x_at	0.003976	gb:NM_002952.1 /DEF=Homo sapiens ribosomal protein S2 (RPS2), mRNA. /FEA=mRNA /GEN=RPS2 /PROD=ribosomal protein S2 /DB_XREF=gi:4506718 /UG=Hs.182426 ribosomal protein S2 /FL=gb:BC001795.1 gb:NM_002952.1	NM_002952		NP_002943
203110_at	0.007174	gb:U43522.1 /DEF=Human cell adhesion kinase beta (CAKbeta) mRNA, complete cds. /FEA=mRNA /GEN=CAKbeta /PROD=cell adhesion kinase beta /DB_XREF=gi:1165218 /UG=Hs.20313 protein tyrosine kinase 2 beta /FL=gb:L49207.1 gb:U43522.1 gb:NM_004103.1 gb:U33284.1	U43522		NP_775268
203113_s_at	7.63E-04	gb:NM_001960.1 /DEF=Homo sapiens eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (EEF1D), mRNA. /FEA=mRNA /GEN=EEF1D /PROD=eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) /DB_XREF=gi:4503478 /UG=Hs.223241 eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) /FL=gb:NM_001960.1	NM_001960		NP_115754

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
203116_s_at	6.73E-04	gb:NM_000140.1 /DEF=Homo sapiens ferrochelatase (protoporphyrin) (FECH), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=FECH /PROD=ferrochelatase precursor /DB_XREF=gi:4557592 /UG=Hs.26 ferrochelatase (protoporphyrin) /FL=gb:NM_000140.1	NM_000140		NP_000131
203120_at	7.63E-04	gb:NM_005426.1 /DEF=Homo sapiens tumor protein p53-binding protein, 2 (TP53BP2), mRNA. /FEA=mRNA /GEN=TP53BP2 /PROD=tumor protein p53-binding protein, 2 /DB_XREF=gi:4885642 /UG=Hs.44585 tumor protein p53-binding protein, 2 /FL=gb:U58334.1 gb:NM_005426.1	NM_005426		NP_005417
203123_s_at	0.025231	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2	AU154469	Hs.57435	NP_000608
203126_at	0.001812	gb:NM_014214.1 /DEF=Homo sapiens inositol(myo)-1(or 4)-monophosphatase 2 (IMPA2), mRNA. /FEA=mRNA /GEN=IMPA2 /PROD=inositol(myo)-1(or 4)-monophosphatase 2 /DB_XREF=gi:7657235 /UG=Hs.5753 inositol(myo)-1(or 4)-monophosphatase 2 /FL=gb:AF200432.1 gb:NM_014214.1 gb:AF014398.2	NM_014214		NP_055029

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203127_s_at	9.85E-04	gb:BC005123.1 /DEF=Homo sapiens, serine palmitoyltransferase, long chain base subunit 2, clone MGC:10362, mRNA, complete cds. /FEA=mRNA /PROD=serine palmitoyltransferase, long chain basesubunit 2 /DB_XREF=gi:13477298 /UG=Hs.59403 serine palmitoyltransferase, long chain base subunit 2 /FL=gb:BC005123.1 gb:AB011098.1 gb:NM_004863.1	BC005123		NP_004854
203128_at	0.043983	gb:NM_004863.1 /DEF=Homo sapiens serine palmitoyltransferase, long chain base subunit 2 (SPTLC2), mRNA. /FEA=mRNA /GEN=SPTLC2 /PROD=serine palmitoyltransferase, long chain basesubunit 2 /DB_XREF=gi:4758667 /UG=Hs.59403 serine palmitoyltransferase, long chain base subunit 2 /FL=gb:BC005123.1 gb:AB011098.1 gb:NM_004863.1	NM_004863		NP_004854
203132_at	0.034296	gb:NM_000321.1 /DEF=Homo sapiens retinoblastoma 1 (including osteosarcoma) (RB1), mRNA. /FEA=mRNA. /GEN=RB1 /PROD=retinoblastoma 1 (including osteosarcoma) /DB_XREF=gi:4506434 /UG=Hs.75770 retinoblastoma 1 (including osteosarcoma) /FL=gb:M33647.1 gb:M15400.1 gb:M28419.1 gb:NM_000321.1	NM_000321		NP_000312

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
203133_at	0.001852	gb:NM_006808.1 /DEF=Homo sapiens protein translocation complex beta (SEC61B), mRNA. /FEA=mRNA /GEN=SEC61B /PROD=protein translocation complex beta /DB_XREF=gi:5803164 /UG=Hs.77028 protein translocation complex beta /FL=gb:BC001734.1 gb:L25085.1 gb:NM_006808.1	NM_006808		NP_006799
203134_at	0.008403	gb:NM_007166.1 /DEF=Homo sapiens Clathrin assembly lymphoid-myeloid leukemia gene (CLTH), mRNA. /FEA=mRNA /GEN=CLTH /PROD=Clathrin assembly lymphoid-myeloid leukemia gene /DB_XREF=gi:6005732 /UG=Hs.7885 phosphatidylinositol binding clathrin assembly protein /FL=gb:U45976.1 gb:NM_007166.1	NM_007166		NP_009097
203137_at	0.014372	gb:NM_004906.1 /DEF=Homo sapiens Wilms tumour 1-associating protein (KIAA0105), mRNA. /FEA=mRNA /GEN=KIAA0105 /PROD=Wilms tumour 1-associating protein /DB_XREF=gi:4758635 /UG=Hs.119 Wilms tumour 1-associating protein /FL=gb:AF277190.1 gb:D14661.1 gb:NM_004906.1	NM_004906		NP_690597

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
203138_at	0.002749	gb:NM_003642.1 /DEF=Homo sapiens histone acetyltransferase 1 (HAT1), mRNA. /FEA=mRNA. /GEN=HAT1 /PROD=histone acetyltransferase 1 /DB_XREF=gi:4504340 /UG=Hs.13340 histone acetyltransferase 1 /FL=gb:AF030424.1 gb:NM_003642.1	NM_003642		NP_003633
203140_at	0.017687	gb:NM_001706.1 /DEF=Homo sapiens B-cell CLLlymphoma 6 (zinc finger protein 51) (BCL6), mRNA. /FEA=mRNA /GEN=BCL6 /PROD=B-cell CLLlymphoma 6 (zinc finger protein 51) /DB_XREF=gi:4502382 /UG=Hs.155024 B-cell CLLlymphoma 6 (zinc finger protein 51) /FL=gb:U00115.1 gb:NM_001706.1	NM_001706		NP_620309
203141_s_at	0.014894	adaptor-related protein complex 3, beta 1 subunit	AW058575	Hs.155172	NP_003655
203142_s_at	7.76E-04	gb:NM_003664.1 /DEF=Homo sapiens adaptor-related protein complex 3, beta 1 subunit (AP3B1), mRNA. /FEA=mRNA /GEN=AP3B1 /PROD=adaptor-related protein complex 3, beta 1 subunit /DB_XREF=gi:4501974 /UG=Hs.155172 adaptor-related protein complex 3, beta 1 subunit /FL=gb:U91931.1 gb:U81504.1 gb:NM_003664.1	NM_003664		NP_003655
203143_s_at	0.014961	KIAA0040 gene product	T79953	Hs.158282	NP_055471

Genes Corresponding To Differentially Expressed Genes in Figur 18 - BladderCancer					
Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203144_s_at	0.025237	gb:NM_014656.1 /DEF=Homo sapiens KIAA0040 gene product (KIAA0040), mRNA. /FEA=mRNA /GEN=KIAA0040 /PROD=KIAA0040 gene product /DB_XREF=gi:7657258 /UG=Hs.158282 KIAA0040 gene product /FL=gb:D25539.1 gb:NM_014656.1	NM_014656		NP_055471
203147_s_at	0.034854	tripartite motif-containing 14	BE962483	Hs.179703	NP_150090
203165_s_at	0.020565	gb:NM_004733.2 /DEF=Homo sapiens acetyl-Coenzyme A transporter (ACATN), mRNA. /FEA=mRNA /GEN=ACATN /PROD=acetyl-Coenzyme A transporter /DB_XREF=gi:6042194 /UG=Hs.285176 acetyl-Coenzyme A transporter /FL=gb:D88152.1 gb:NM_004733.2	NM_004733		NP_004724
203166_at	0.027219	gb:NM_006324.1 /DEF=Homo sapiens craniofacial development protein 1 (CFDP1), mRNA. /FEA=mRNA /GEN=CFDP1 /PROD=craniofacial development protein 1 /DB_XREF=gi:5453566 /UG=Hs.296460 craniofacial development protein 1 /FL=gb:BC000991.2 gb:AB009285.1 gb:NM_006324.1	NM_006324		NP_006315
203167_at	7.63E-04	gb:NM_003255.2 /DEF=Homo sapiens tissue inhibitor of metalloproteinase 2 (TIMP2), mRNA. /FEA=mRNA /GEN=TIMP2 /PROD=tissue inhibitor of metalloproteinase 2precursor /DB_XREF=gi:9257247 /UG=Hs.325495 tissue inhibitor of metalloproteinase 2 /FL=gb:M32304.1 gb:J05593.1 gb:NM_003255.2	NM_003255		NP_003246

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203184_at	0.001719	gb:NM_001999.2 /DEF=Homo sapiens fibrillin 2 (congenital contractural arachnodactyly) (FBN2), mRNA. /FEA=mRNA /GEN=FBN2 /PROD=fibrillin 2 /DB_XREF=gi:4755135 /UG=Hs.79432 fibrillin 2 (congenital contractural arachnodactyly) /FL=gb:U03272.1 gb:NM_001999.2	NM_001999		NP_001990
203185_at	0.007503	gb:NM_014737.1 /DEF=Homo sapiens Ras association (RalGDSAF-6) domain family 2 (RASSF2), mRNA. /FEA=mRNA /GEN=RASSF2 /PROD=Ras association (RalGDSAF-6) domain family 2 /DB_XREF=gi:7661963 /UG=Hs.80905 Ras association (RalGDSAF-6) domain family 2 /FL=gb:D79990.1 gb:NM_014737.1	NM_014737		NP_739580
203186_s_at	0.009378	gb:NM_002961.2 /DEF=Homo sapiens S100 calcium-binding protein A4 (calcium protein, calvasculin, metastasin, murine placental homolog) (S100A4), transcript variant 1, mRNA. /FEA=mRNA /GEN=S100A4 /PROD=S100 calcium-binding protein A4 /DB_XREF=gi:9845514 /UG=Hs.81256 S100 calcium-binding protein A4 (calcium protein, calvasculin, metastasin, murine placental homolog) /FL=gb:NM_002961.2 gb:NM_019554.1	NM_002961		NP_062427
203196_at	0.033831	ATP-binding cassette, sub-family C (CFTR/MRP), member 4	AI948503	Hs.387834	NP_005836
203197_s_at	0.038535	carnitine palmitoyltransferase II	AW157077	Hs.274336	NP_060357
203202_at	0.004519	glioma pathogenesis-related protein	AI950314	Hs.154762	NP_008974

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203203_s_at	0.001286	gb:NM_007043.1 /DEF=Homo sapiens HIV-1 rev binding protein 2 (HRB2), mRNA. /FEA=mRNA /GEN=HRB2 /PROD=HIV-1 rev binding protein 2 /DB_XREF=gi:5902047 /UG=Hs.154762 HIV-1 rev binding protein 2 /FL=gb:U55766.1 gb:NM_007043.1	NM_007043		NP_008974
203204_s_at	0.020272	gb:BC002558.1 /DEF=Homo sapiens, KIAA0677 gene product, clone MGC:1972, mRNA, complete cds. /FEA=mRNA /PROD=KIAA0677 gene product /DB_XREF=gi:12803466 /UG=Hs.155983 KIAA0677 gene product /FL=gb:BC002558.1 gb:AB014577.1 gb:NM_014663.1	BC002558		NP_055478
203206_at	0.002749	gb:NM_014661.1 /DEF=Homo sapiens KIAA0140 gene product (KIAA0140), mRNA. /FEA=mRNA /GEN=KIAA0140 /PROD=KIAA0140 gene product /DB_XREF=gi:7661937 /UG=Hs.156016 KIAA0140 gene product /FL=gb:D50930.1 gb:NM_014661.1	NM_014661		NP_055476
203213_at	0.01148	cell division cycle 2, G1 to S and G2 to M	AL524035	Hs.334562	NP_203698
203223_at	0.034317	gb:NM_004703.1 /DEF=Homo sapiens rabaptin-5 (RAB5EP), mRNA. /FEA=mRNA /GEN=RAB5EP /PROD=rabaptin-5 /DB_XREF=gi:4759005 /UG=Hs.326056 rabaptin-5 /FL=gb:NM_004703.1	NM_004703		NP_004694

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203232_s_at	0.00121	gb:NM_000332.1 /DEF=Homo sapiens spinocerebellar ataxia 1 (olivopontocerebellar ataxia 1, autosomal dominant, ataxin 1) (SCA1), mRNA. /FEA=mRNA /GEN=SCA1 /PROD=ataxin 1 /DB_XREF=gi:4506792 /UG=Hs.74520 spinocerebellar ataxia 1 (olivopontocerebellar ataxia 1, autosomal dominant, ataxin 1) /FL=gb:NM_000332.1	NM_000332		
203234_at	0.003564	gb:NM_003364.1 /DEF=Homo sapiens uridine phosphorylase (UP), mRNA. /FEA=mRNA /GEN=UP /PROD=uridine phosphorylase /DB_XREF=gi:4507838 /UG=Hs.77573 uridine phosphorylase /FL=gb:BC001405.1 gb:NM_003364.1	NM_003364		NP_003355
203243_s_at	0.001763	gb:NM_006457.1 /DEF=Homo sapiens LIM protein (similar to rat protein kinase C-binding enigma) (LIM), mRNA. /FEA=mRNA /GEN=LIM /PROD=LIM protein (similar to rat protein kinase C-binding enigma) /DB_XREF=gi:5453713 /UG=Hs.154103 LIM protein (similar to rat protein kinase C-binding enigma) /FL=gb:AF061258.1 gb:NM_006457.1	NM_006457		NP_006448
203244_at	0.014472	gb:NM_000319.1 /DEF=Homo sapiens peroxisome receptor 1 (PXR1), mRNA. /FEA=mRNA /GEN=PXR1 /PROD=peroxisome receptor 1 /DB_XREF=gi:4506346 /UG=Hs.158084 peroxisome receptor 1 /FL=gb:NM_000319.1 gb:U19721.1	NM_000319		NP_000310

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203247_s_at	0.006756	gb:BC003566.1 /DEF=Homo sapiens, zinc finger protein 24 (KOX 17), clone. MGC:2057, mRNA, complete cds. /FEA=mRNA /PROD=zinc finger protein 24 (KOX 17) /DB_XREF=gi:13097725 /UG=Hs.183593 zinc finger protein 24 (KOX 17) /FL=gb:BC003566.1 gb:U68536.1 gb:AF038964.1 gb:NM_006965.1	BC003566		NP_008896
203253_s_at	0.004519	gb:NM_015216.1 /DEF=Homo sapiens KIAA0433 protein (KIAA0433), mRNA. /FEA=mRNA /GEN=KIAA0433 /PROD=KIAA0433 protein /DB_XREF=gi:7662117 /UG=Hs.26179 KIAA0433 protein /FL=gb:NM_015216.1	NM_015216		NP_056031
203265_s_at	7.76E-04	mitogen-activated protein kinase kinase 4	AA810268	Hs.75217	
203266_s_at	0.001213	gb:NM_003010.1 /DEF=Homo sapiens mitogen-activated protein kinase kinase 4 (MAP2K4), mRNA. /FEA=mRNA /GEN=MAP2K4 /PROD=mitogen-activated protein kinase kinase 4 /DB_XREF=gi:4506888 /UG=Hs.75217 mitogen-activated protein kinase kinase 4 /FL=gb:NM_003010.1 gb:L36870.1 gb:U17743.1	NM_003010		NP_003001
203272_s_at	0.020839	lung cancer candidate	BF308548	Hs.8186	NP_009206
203273_s_at	0.015018	gb:NM_007275.1 /DEF=Homo sapiens lung cancer candidate (FUS1), mRNA. /FEA=mRNA /GEN=FUS1 /PROD=lung cancer candidate /DB_XREF=gi:6005759 /UG=Hs.8186 lung cancer candidate /FL=gb:AF055479.1 gb:NM_007275.1	NM_007275		NP_009206

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203274_at	6.66E-04	gb:NM_012151.2 /DEF=Homo sapiens coagulation factor VIII-associated (intronic transcript) (F8A), mRNA. /FEA=mRNA /GEN=F8A /PROD=coagulation factor VIII-associated protein /DB_XREF=gi:12056462 /UG=Hs.83363 coagulation factor VIII-associated (intronic transcript) /FL=gb:NM_012151.2	NM_012151		NP_036283
203275_at	0.00643	gb:NM_002199.2 /DEF=Homo sapiens interferon regulatory factor 2 (IRF2), mRNA. /FEA=mRNA /GEN=IRF2 /PROD=interferon regulatory factor 2 /DB_XREF=gi:4755144 /UG=Hs.83795 interferon regulatory factor 2 /FL=gb:NM_002199.2	NM_002199		NP_002190
203279_at	0.001763	gb:NM_014674.1 /DEF=Homo sapiens KIAA0212 gene product (KIAA0212), mRNA. /FEA=mRNA /GEN=KIAA0212 /PROD=KIAA0212 gene product /DB_XREF=gi:7662001 /UG=Hs.154332 KIAA0212 gene product /FL=gb:D86967.1 gb:NM_014674.1	NM_014674		NP_055489

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203282_at	6.66E-04	gb:NM_000158.1 /DEF=Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA. /FEA=mRNA /GEN=GBE1 /PROD=glucan (1,4-alpha-), branching enzyme 1(glycogen branching enzyme) /DB_XREF=gi:4557618 /UG=Hs.1691 glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) /FL=gb:L07956.1 gb:NM_000158.1	NM_000158		NP_000149
203283_s_at	0.041166	Consensus includes gb:AK023260.1 /DEF=Homo sapiens cDNA FLJ13198 fis, clone NT2RP3004454, highly similar to Homo sapiens mRNA for KIAA0448 protein. /FEA=mRNA /DB_XREF=gi:10435114 /UG=Hs.169939 heparan sulfate 2-O-sulfotransferase /FL=gb:NM_012262.2 gb:AB007917.1 gb:AB024568.1	NM_012262		NP_036394
203284_s_at	0.001852	Consensus includes gb:AW151887 /FEA=EST /DB_XREF=gi:6199872 /DB_XREF=est:xf72c12.x1 /CLONE=IMAGE:2623606 /UG=Hs.169939 heparan sulfate 2-O-sulfotransferase /FL=gb:NM_012262.2 gb:AB007917.1 gb:AB024568.1	NM_012262		NP_036394

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203285_s_at	7.25E-04	gb:NM_012262.2 /DEF=Homo sapiens heparan sulfate 2-O-sulfotransferase (HS2ST1), mRNA. /FEA=mRNA /GEN=HS2ST1 /PROD=heparan sulfate 2-O-sulfotransferase 1 /DB_XREF=gi:12545388 /UG=Hs.169939 heparan sulfate 2-O-sulfotransferase /FL=gb:NM_012262.2 gb:AB007917.1 gb:AB024568.1	NM_012262		NP_036394
203286_at	9.85E-04	gb:NM_014901.1 /DEF=Homo sapiens KIAA1100 protein (KIAA1100), mRNA. /FEA=mRNA /GEN=KIAA1100 /PROD=KIAA1100 protein /DB_XREF=gi:7662485 /UG=Hs.179946 KIAA1100 protein /FL=gb:AB029023.1 gb:NM_014901.1	NM_014901		NP_055716
203288_at	0.015899	gb:NM_014686.1 /DEF=Homo sapiens KIAA0355 gene product (KIAA0355), mRNA. /FEA=mRNA /GEN=KIAA0355 /PROD=KIAA0355 gene product /DB_XREF=gi:7662075 /UG=Hs.186840 KIAA0355 gene product /FL=gb:AB002353.1 gb:NM_014686.1	NM_014686		NP_055501
203291_at	0.002446	gb:NM_013316.1 /DEF=Homo sapiens CCR4-NOT transcription complex, subunit 4 (CNOT4), mRNA. /FEA=mRNA /GEN=CNOT4 /PROD=CCR4-NOT transcription complex, subunit 4 /DB_XREF=gi:7019466 /UG=Hs.20423 CCR4-NOT transcription complex, subunit 4 /FL=gb:U71267.1 gb:NM_013316.1	NM_013316		NP_037448

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
203292_s_at	0.005347	gb:NM_021729.2 /DEF=Homo sapiens vacuolar protein sorting 11 (yeast homolog) (VPS11), mRNA. /FEA=mRNA /GEN=VPS11 /PROD=vacuolar protein sorting 11 (yeast homolog) /DB_XREF=gi:11141864 /UG=Hs.234282 vacuolar protein sorting 11 (yeast homolog) /FL=gb:AB027508.2 gb:NM_021729.2 gb:AF308800.1	NM_021729		NP_068375
203300_x_at	0.047167	gb:NM_003916.1 /DEF=Homo sapiens adaptor-related protein complex 1, sigma 2 subunit (AP1S2), mRNA. /FEA=mRNA /GEN=AP1S2 /PROD=adaptor-related protein complex 1, sigma 2 subunit /DB_XREF=gi:4506956 /UG=Hs.40368 adaptor-related protein complex 1, sigma 2 subunit /FL=gb:AF251295.1 gb:BC001117.1 gb:AB015320.1 gb:NM_003916.1	NM_003916		NP_003907
203302_at	0.021589	gb:NM_000788.1 /DEF=Homo sapiens deoxycytidine kinase (DCK), mRNA. /FEA=mRNA /GEN=DCK /PROD=deoxycytidine kinase /DB_XREF=gi:4503268 /UG=Hs.709 deoxycytidine kinase /FL=gb:M60527.1 gb:NM_000788.1	NM_000788		NP_000779
203304_at	0.029244	gb:NM_012342.1 /DEF=Homo sapiens putative transmembrane protein (NMA), mRNA. /FEA=mRNA /GEN=NMA /PROD=putative transmembrane protein /DB_XREF=gi:6912533 /UG=Hs.78776 putative transmembrane protein /FL=gb:U23070.1 gb:NM_012342.1	NM_012342		NP_036474

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
203305_at	0.041179	gb:NM_000129.2 /DEF=Homo sapiens coagulation factor XIII, A1 polypeptide (F13A1), mRNA. /FEA=mRNA /GEN=F13A1 /PROD=coagulation factor XIII A1 subunit precursor /DB_XREF=gi:9961355 /UG=Hs.80424 coagulation factor XIII, A1 polypeptide. /FL=gb:M14354.1 gb:NM_000129.2	NM_000129		NP_000120
203306_s_at	0.032443	gb:NM_006416.1 /DEF=Homo sapiens solute carrier family 35 (CMP-sialic acid transporter), member 1 (SLC35A1), mRNA. /FEA=mRNA /GEN=SLC35A1 /PROD=solute carrier family 35 (CMP-sialic acid transporter), member 1 /DB_XREF=gi:5453620 /UG=Hs.82921 solute carrier family 35 (CMP-sialic acid transporter), member 1 /FL=gb:D87969.1 gb:NM_006416.1	NM_006416		NP_006407
203309_s_at	0.006483	gb:NM_000195.1 /DEF=Homo sapiens Hermansky-Pudlak syndrome (HPS), mRNA. /FEA=mRNA /GEN=HPS /PROD=Hermansky-Pudlak syndrome protein /DB_XREF=gi:4504484 /UG=Hs.83951 Hermansky-Pudlak syndrome /FL=gb:U65676.1 gb:NM_000195.1	NM_000195		NP_000186

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203310_at	0.018534	gb:NM_007269.1 /DEF=Homo sapiens syntaxin binding protein 3 (STXBP3), mRNA. /FEA=mRNA /GEN=STXBP3 /PROD=syntaxin 4 binding protein /DB_XREF=gi:6005885 /UG=Hs.8813 syntaxin binding protein 3 /FL=gb:D63506.1 gb:AF032922.1 gb:NM_007269.1	NM_007269		NP_009200
203318_s_at	7.76E-04	gb:NM_021964.1 /DEF=Homo sapiens zinc finger protein 148 (pHZ-52) (ZNF148), mRNA. /FEA=mRNA /GEN=ZNF148 /PROD=zinc finger protein 148 (pHZ-52) /DB_XREF=gi:11415035 /UG=Hs.112180 zinc finger protein 148 (pHZ-52) /FL=gb:NM_021964.1 gb:L04282.1	NM_021964		NP_068799
203319_s_at	0.003875	gb:L04282.1 /DEF=Human CACCC box-binding protein mRNA, complete cds. /FEA=mRNA /PROD=CACCC box-binding protein /DB_XREF=gi:388318 /UG=Hs.112180 zinc finger protein 148 (pHZ-52) /FL=gb:NM_021964.1 gb:L04282.1	L04282		NP_068799
203320_at	0.001763	gb:NM_005475.1 /DEF=Homo sapiens lymphocyte adaptor protein (LNK), mRNA. /FEA=mRNA /GEN=LNK /PROD=lymphocyte adaptor protein /DB_XREF=gi:4885454 /UG=Hs.13131 lymphocyte adaptor protein /FL=gb:AF055581.1 gb:NM_005475.1	NM_005475		NP_005466

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203321_s_at	0.001719	Consensus includes gb:AK022688.1 /DEF=Homo sapiens cDNA FLJ12626 fis, clone NT2RM4001810, weakly similar to AGGRECAN CORE PROTEIN PRECURSOR. /FEA=CDS /DB_XREF=gi:10434228 /UG=Hs.131915 KIAA0863 protein /FL=gb:AB020670.1 gb:NM_014913.1	NM_014913		NP_055728
203328_x_at	0.048713	gb:NM_004969.1 /DEF=Homo sapiens insulin-degrading enzyme (IDE), mRNA. /FEA=mRNA /GEN=IDE /PROD=insulysin /DB_XREF=gi:4826769 /UG=Hs.1508 insulin-degrading enzyme /FL=gb:M21188.1 gb:NM_004969.1	NM_004969		NP_004960
203340_s_at	0.032979	gb:NM_003705.1 /DEF=Homo sapiens solute carrier family 25 (mitochondrial carrier, Aralar), member 12 (SLC25A12), mRNA. /FEA=mRNA /GEN=SLC25A12 /PROD=solute carrier family 25 (mitochondrial carrier, Aralar), member 12 /DB_XREF=gi:4507006 /UG=Hs.179866 solute carrier family 25 (mitochondrial carrier, Aralar), member 12 /FL=gb:NM_003705.1	NM_003705		NP_003696

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203342_at	0.016588	gb:NM_005834.1 /DEF=Homo sapiens translocase of inner mitochondrial membrane 17 (yeast) homolog B (TIM17B), mRNA. /FEA=mRNA /GEN=TIM17B /PROD=translocase of inner mitochondrial membrane 17 (yeast) homolog B /DB_XREF=gi:5032180 /UG=Hs.19105 translocase of inner mitochondrial membrane 17 (yeast) homolog B /FL=gb:AJ005895.1 gb:AF034790.1 gb:AF077039.1 gb:NM_005834.1	NM_005834		NP_005825
203347_s_at	0.023393	gb:NM_007358.1 /DEF=Homo sapiens putative DNA binding protein (M96), mRNA. /FEA=mRNA /GEN=M96 /PROD=putative DNA binding protein /DB_XREF=gi:6678763 /UG=Hs.31016 putative DNA binding protein /FL=gb:AF072814.1 gb:NM_007358.1	NM_007358		NP_031384
203353_s_at	0.020993	gb:NM_015846.1 /DEF=Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant 1, mRNA. /FEA=mRNA /GEN=MBD1 /PROD=methyl-CpG binding domain protein 1, isoform 1 /DB_XREF=gi:7710138 /UG=Hs.6211 methyl-CpG binding domain protein 1 /FL=gb:AF078830.1 gb:NM_015846.1	NM_015846		NP_056723
203359_s_at	6.66E-04	c-myc binding protein	AL525412	Hs.78221	NP_036465

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203360_s_at	6.66E-04	gb:D50692.1 /DEF=Homo sapiens mRNA for c-myc binding protein, complete cds. /FEA=mRNA /PROD=c-myc binding protein /DB_XREF=gi:1785850 /UG=Hs.78221 c-myc binding protein /FL=gb:D50692.1 gb:AB007191.2 gb:NM_012333.2	D50692		NP_036465
203361_s_at	0.038584	gb:NM_012333.2 /DEF=Homo sapiens c-myc binding protein (MYCBP), mRNA. /FEA=mRNA /GEN=MYCBP /PROD=c-myc binding protein /DB_XREF=gi:8850230 /UG=Hs.78221 c-myc binding protein /FL=gb:D50692.1 gb:AB007191.2 gb:NM_012333.2	NM_012333		NP_036465
203362_s_at	0.002931	gb:NM_002358.2 /DEF=Homo sapiens MAD2 (mitotic arrest deficient, yeast, homolog)-like 1 (MAD2L1), mRNA. /FEA=mRNA /GEN=MAD2L1 /PROD=MAD2-like 1 /DB_XREF=gi:6466452 /UG=Hs.79078 MAD2 (mitotic arrest deficient, yeast, homolog)-like 1 /FL=gb:BC000356.1 gb:U65410.1 gb:NM_002358.2 gb:U31278.1	NM_002358		NP_002349

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203371_s_at	0.002603	gb:NM_002491.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, B12) (NDUFB3), mRNA. /FEA=mRNA /GEN=NDUFB3 /PROD=NADH dehydrogenase (ubiquinone) 1 betasubcomplex, 3 (12kD, B12) /DB_XREF=gi:4505360 /UG=Hs.109760 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, B12) /FL=gb:AF047183.1 gb:NM_002491.1	NM_002491		NP_002482
203376_at	0.030752	pre-mRNA splicing factor 17	BG528818	Hs.116674	NP_056975
203379_at	0.002724	gb:NM_002953.1 /DEF=Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 1 (RPS6KA1), mRNA. /FEA=mRNA /GEN=RPS6KA1 /PROD=ribosomal protein S6 kinase, 90kD, polypeptide1 /DB_XREF=gi:4506732 /UG=Hs.149957 ribosomal protein S6 kinase, 90kD, polypeptide 1. /FL=gb:L07597.1 gb:NM_002953.1	NM_002953		NP_002944
203383_s_at	0.039351	golgi autoantigen, golgin subfamily a, 1	BG111661	Hs.172647	NP_002068
203384_s_at	0.024403	gb:NM_002077.1 /DEF=Homo sapiens golgi autoantigen, golgin subfamily a, 1 (GOLGA1), mRNA. /FEA=mRNA /GEN=GOLGA1 /PROD=golgin 97 /DB_XREF=gi:4504062 /UG=Hs.172647 golgi autoantigen, golgin subfamily a, 1 /FL=gb:U51587.1 gb:NM_002077.1	NM_002077		NP_002068

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
203385_at	0.024696	gb:NM_001345.1 /DEF=Homo sapiens diacylglycerol kinase, alpha (80kD) (DGKA), mRNA. /FEA=mRNA /GEN=DGKA /PROD=diacylglycerol kinase, alpha (80kD) /DB_XREF=gi:11415023 /UG=Hs.172690 diacylglycerol kinase, alpha (80kD) /FL=gb:NM_001345.1 gb:AF064770.1	NM_001345		NP_001336
203387_s_at	0.044364	gb:NM_014832.1 /DEF=Homo sapiens KIAA0603 gene product (KIAA0603), mRNA. /FEA=mRNA /GEN=KIAA0603 /PROD=KIAA0603 gene product /DB_XREF=gi:7662197 /UG=Hs.173802 KIAA0603 gene product /FL=gb:AB011175.1 gb:NM_014832.1	NM_014832		NP_055647
203394_s_at	0.009582	hairy homolog (Drosophila)	BE973687	Hs.250666	NP_005515
203397_s_at	0.014263	UDP-N-acetyl-alpha-D-galactosamine:polypeptide-N-acetylgalactosaminyltransferase 3 (GalNAc-T3)	BF063271	Hs.278611	NP_004473
203406_at	0.011135	gb:NM_005926.1 /DEF=Homo sapiens microfibrillar-associated protein 1 (MFAP1), mRNA. /FEA=mRNA /GEN=MFAP1 /PROD=microfibrillar-associated protein 1 /DB_XREF=gi:5174552 /UG=Hs.61418 microfibrillar-associated protein 1 /FL=gb:U04209.1 gb:NM_005926.1	NM_005926		NP_005917

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203408_s_at	0.001852	gb:NM_002971.1 /DEF=Homo sapiens special AT-rich sequence binding protein 1 (binds to nuclear matrixscaffold-associating DNAs) (SATB1), mRNA. /FEA=mRNA /GEN=SATB1 /PROD=special AT-rich sequence binding protein 1(binds to nuclear matrixscaffold-associating DNAs) /DB_XREF=gi:4506790 /UG=Hs.74592 special AT-rich sequence binding protein 1 (binds to nuclear matrixscaffold-associating DNAs) /FL=gb:M97287.1 gb:NM_002971.1	NM_002971		NP_002962
203415_at	7.63E-04	gb:NM_013232.1 /DEF=Homo sapiens programmed cell death 6 (PDCD6), mRNA. /FEA=mRNA /GEN=PDCD6 /PROD=programmed cell death 6 /DB_XREF=gi:7019484 /UG=Hs.80019 programmed cell death 6 /FL=gb:AF035606.1 gb:U58773.1 gb:NM_013232.1	NM_013232		NP_037364
203416_at	0.00643	gb:NM_000560.1 /DEF=Homo sapiens CD53 antigen (CD53), mRNA. /FEA=mRNA /GEN=CD53 /PROD=CD53 antigen /DB_XREF=gi:10834971 /UG=Hs.82212 CD53 antigen /FL=gb:NM_000560.1 gb:M60871.1 gb:M37033.1	NM_000560		NP_000551
203430_at	6.66E-04	gb:NM_014320.1 /DEF=Homo sapiens putative heme-binding protein (SOUL), mRNA. /FEA=mRNA /GEN=SOUL /PROD=putative heme-binding protein /DB_XREF=gi:7657602 /UG=Hs.111029 putative heme-binding protein /FL=gb:AF117616.1 gb:NM_014320.1	NM_014320		NP_055135

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203432_at	0.044397	thymopoietin	AW272611	Hs.11355	NP_003267
203433_at	0.004311	gb:NM_006441.1 /DEF=Homo sapiens 5,10-methenyltetrahydrofolate synthetase (5-formyltetrahydrofolate cyclo-ligase) (MTHFS), mRNA. /FEA=mRNA /GEN=MTHFS /PROD=5,10-methenyltetrahydrofolate synthetase(5-formyltetrahydrofolate cyclo-ligase) /DB_XREF=gi:5453745 /UG=Hs.118131 5,10-methenyltetrahydrofolate synthetase (5-formyltetrahydrofolate cyclo-ligase) /FL=gb:NM_006441.1 gb:L38928.1	NM_006441		NP_006432
203435_s_at	0.043386	gb:NM_007287.1 /DEF=Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) (MME), transcript variant 1bis, mRNA. /FEA=mRNA /GEN=MME /PROD=membrane metallo-endopeptidase /DB_XREF=gi:6042199 /UG=Hs.1298 membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) /FL=gb:J03779.1 gb:NM_007287.1 gb:NM_007288.1	NM_007287		NP_009220
203437_at	7.63E-04	gb:NM_003876.1 /DEF=Homo sapiens putative receptor protein (PMI), mRNA. /FEA=mRNA /GEN=PMI /PROD=putative receptor protein /DB_XREF=gi:4505900 /UG=Hs.15196 putative receptor protein /FL=gb:BC002819.1 gb:BC005268.1 gb:NM_003876.1	NM_003876		NP_003867

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203445_s_at	0.006596	gb:NM_005730.1 /DEF=Homo sapiens conserved gene amplified in osteosarcoma (OS4), mRNA. /FEA=mRNA /GEN=OS4 /PROD=conserved gene amplified in osteosarcoma /DB_XREF=gi:5031964 /UG=Hs.180669 conserved gene amplified in osteosarcoma /FL=gb:U81556.1 gb:NM_005730.1	NM_005730		NP_005721
203447_at	0.032443	proteasome (prosome, macropain) 26S subunit, non-ATPase, 5	AU157008	Hs.273721	NP_005038
203457_at	0.00121	gb:NM_003569.1 /DEF=Homo sapiens syntaxin 7 (STX7), mRNA. /FEA=mRNA /GEN=STX7 /PROD=syntaxin 7 /DB_XREF=gi:4507294 /UG=Hs.299166 syntaxin 7 /FL=gb:U77942.1 gb:NM_003569.1	NM_003569		NP_003560
203460_s_at	7.25E-04	gb:NM_007318.1 /DEF=Homo sapiens presenilin 1 (Alzheimer disease 3) (PSEN1), transcript variant I-463, mRNA. /FEA=mRNA /GEN=PSEN1 /PROD=presenilin 1 isoform I-463 /DB_XREF=gi:7549812 /UG=Hs.3260 presenilin 1 (Alzheimer disease 3) /FL=gb:U40379.1 gb:L76517.1 gb:NM_007318.1	NM_007318		NP_015558
203471_s_at	0.002603	gb:NM_002664.1 /DEF=Homo sapiens pleckstrin (PLEK), mRNA. /FEA=mRNA /GEN=PLEK /PROD=pleckstrin /DB_XREF=gi:4505878 /UG=Hs.77436 pleckstrin /FL=gb:NM_002664.1	NM_002664		NP_002655

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203478_at	0.018947	gb:NM_002494.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (6kD, KFYI) (NDUFC1), mRNA. /FEA=mRNA /GEN=NDUFC1 /PROD=NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (6kD, KFYI) /DB_XREF=gi:4505366 /UG=Hs.84549 NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (6kD, KFYI) /FL=gb:AF047184.1 gb:AF047435.1 gb:NM_002494.1	NM_002494		NP_002485
203484_at	0.018901	gb:NM_014302.1 /DEF=Homo sapiens Sec61 gamma (SEC61G), mRNA. /FEA=mRNA /GEN=SEC61G /PROD=Sec61 gamma /DB_XREF=gi:7657545 /UG=Hs.9950 Sec61 gamma /FL=gb:AF054184.1 gb:NM_014302.1	NM_014302		NP_055117
203486_s_at	0.001008	DKFZP434A043 protein	BF195973	Hs.102708	NP_056211
203487_s_at	0.001763	gb:NM_015396.1 /DEF=Homo sapiens DKFZP434A043 protein (DKFZP434A043), mRNA. /FEA=mRNA /GEN=DKFZP434A043 /PROD=DKFZP434A043 protein /DB_XREF=gi:7661561 /UG=Hs.102708 DKFZP434A043 protein /FL=gb:NM_015396.1	NM_015396		NP_056211

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203497_at	0.002446	gb:NM_004774.1 /DEF=Homo sapiens PPAR binding protein (PPARBP), mRNA. /FEA=mRNA /GEN=PPARBP /PROD=thyroid hormone receptor interactor 2 /DB_XREF=gi:4759265 /UG=Hs.15589 PPAR binding protein /FL=gb:AF055994.1 gb:NM_004774.1 gb:AF283812.1	NM_004774		NP_004765
203501_at	0.001719	gb:NM_006102.1 /DEF=Homo sapiens plasma glutamate carboxypeptidase (PGCP), mRNA. /FEA=mRNA /GEN=PGCP /PROD=plasma glutamate carboxypeptidase /DB_XREF=gi:5174626 /UG=Hs.197335 plasma glutamate carboxypeptidase /FL=gb:AF119386.1 gb:NM_006102.1	NM_006102		
203502_at	6.66E-04	gb:NM_001724.1 /DEF=Homo sapiens 2,3-bisphosphoglycerate mutase (BPGM), mRNA. /FEA=mRNA /GEN=BPGM /PROD=2,3-bisphosphoglycerate mutase /DB_XREF=gi:4502444 /UG=Hs.198365 2,3-bisphosphoglycerate mutase /FL=gb:NM_001724.1	NM_001724		NP_001715

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203506_s_at	0.003315	gb:NM_005120.1 /DEF=Homo sapiens trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit) (TNRC11), mRNA. /FEA=mRNA /GEN=TNRC11 /PROD=trinucleotide repeat containing 11(THR-associated protein, 230 kDa subunit) /DB_XREF=gi:4827041 /UG=Hs.211607 trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit) /FL=gb:U80742.1 gb:AF071309.1 gb:AF117755.1 gb:NM_005120.1	NM_005120		NP_005111
203507_at	0.002536	gb:NM_001251.1 /DEF=Homo sapiens CD68 antigen (CD68), mRNA. /FEA=mRNA /GEN=CD68 /PROD=CD68 antigen /DB_XREF=gi:4557434 /UG=Hs.246381 CD68 antigen /FL=gb:NM_001251.1	NM_001251		NP_001242
203508_at	0.00109	gb:NM_001066.1 /DEF=Homo sapiens tumor necrosis factor receptor superfamily, member 1B (TNFRSF1B), mRNA. /FEA=mRNA /GEN=TNFRSF1B /PROD=tumor necrosis factor receptor 2 (75kD) /DB_XREF=gi:4507576 /UG=Hs.256278 tumor necrosis factor receptor superfamily, member 1B /FL=gb:M32315.1 gb:M55994.1 gb:NM_001066.1	NM_001066		NP_001057
203511_s_at	0.002749	gb:AF041432.1 /DEF=Homo sapiens bet3 (BET3) mRNA, complete cds. /FEA=mRNA /GEN=BET3 /PROD=bet3 /DB_XREF=gi:2791803 /UG=Hs.288013 similar to yeast BET3 (S. cerevisiae) /FL=gb:AF041432.1 gb:NM_014408.1	AF041432		NP_055223

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203512_at	0.002603	gb:NM_014408.1 /DEF=Homo sapiens similar to yeast BET3 (S. cerevisiae) (BET3), mRNA. /FEA=mRNA /GEN=BET3 /PROD=similar to yeast BET3 (S. cerevisiae) /DB_XREF=gi:7656925 /UG=Hs.288013 similar to yeast BET3 (S. cerevisiae) /FL=gb:AF041432.1 gb:NM_014408.1	NM_014408		NP_055223
203513_at	0.030968	gb:NM_025137.1 /DEF=Homo sapiens hypothetical protein FLJ21439 (FLJ21439), mRNA. /FEA=mRNA /GEN=FLJ21439 /PROD=hypothetical protein FLJ21439 /DB_XREF=gi:13376718 /UG=Hs.288872 hypothetical protein FLJ21439 /FL=gb:NM_025137.1	NM_025137		NP_079413
203514_at	0.03867	mitogen-activated protein kinase kinase kinase 3	BF971923	Hs.29282	NP_002392
203518_at	0.004987	gb:NM_000081.1 /DEF=Homo sapiens Chediak-Higashi syndrome 1 (CHS1), mRNA. /FEA=mRNA /GEN=CHS1 /PROD=beige protein homolog /DB_XREF=gi:4502838 /UG=Hs.36508 Chediak-Higashi syndrome 1. /FL=gb:U67615.1 gb:NM_000081.1	NM_000081		NP_000072
203522_at	0.038994	gb:NM_005125.1 /DEF=Homo sapiens copper chaperone for superoxide dismutase (CCS), mRNA. /FEA=mRNA /GEN=CCS /PROD=copper chaperone for superoxide dismutase /DB_XREF=gi:4826664 /UG=Hs.5002 copper chaperone for superoxide dismutase /FL=gb:AF002210.1 gb:NM_005125.1	NM_005125		NP_005116

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203525_s_at	0.024403	adenomatosis polyposis coli	AI375486	Hs.75081	NP_000029
203529_at	0.002603	gb:NM_016294.1 /DEF=Homo sapiens serinethreonine protein phosphatase catalytic subunit (LOC51723), mRNA. /FEA=mRNA /GEN=LOC51723 /PROD=serinethreonine protein phosphatase catalytic subunit /DB_XREF=gi:7706488 /UG=Hs.80324 serinethreonine protein phosphatase catalytic subunit /FL=gb:AF035158.2 gb:NM_016294.1	NM_016294		
203534_at	0.011969	gb:NM_014462.1 /DEF=Homo sapiens Lsm1 protein (LSM1), mRNA. /FEA=mRNA /GEN=LSM1 /PROD=Lsm1 protein /DB_XREF=gi:7657312 /UG=Hs.111783 Lsm1 protein /FL=gb:BC001767.1 gb:AF000177.1 gb:NM_014462.1	NM_014462		NP_055277
203544_s_at	0.001763	gb:NM_003473.1 /DEF=Homo sapiens signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 (STAM), mRNA. /FEA=mRNA /GEN=STAM /PROD=signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 /DB_XREF=gi:4507248 /UG=Hs.153487 signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 /FL=gb:U43899.1 gb:NM_003473.1	NM_003473		NP_003464

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203553_s_at	0.009935	gb:NM_006575.1 /DEF=Homo sapiens mitogen-activated protein kinase kinase kinase kinase 5 (MAP4K5), mRNA. /FEA=mRNA /GEN=MAP4K5 /PROD=mitogen-activated protein kinase kinase kinasekinase /DB_XREF=gi:5729890 /UG=Hs.246970 mitogen-activated protein kinase kinase kinase 5. /FL=gb:U77129.1 gb:NM_006575.1	NM_006575		NP_006566
203560_at	7.60E-04	gb:NM_003878.1 /DEF=Homo sapiens gamma-glutamyl hydrolase (conjugase, folylpolygammaglutamyl hydrolase) (GGH), mRNA. /FEA=mRNA /GEN=GGH /PROD=gamma-glutamyl hydrolase (conjugase,folypolygammaglutamyl hydrolase) precursor /DB_XREF=gi:4503986 /UG=Hs.78619 gamma-glutamyl hydrolase (conjugase, folypolygammaglutamyl hydrolase) /FL=gb:U55206.1 gb:NM_003878.1	NM_003878		NP_003869
203561_at	0.005347	gb:NM_021642.1 /DEF=Homo sapiens Fc fragment of IgG, low affinity IIa, receptor for (CD32) (FCGR2A), mRNA. /FEA=mRNA /GEN=FCGR2A /PROD=Fc fragment of IgG, low affinity IIa, receptorfor (CD32) /DB_XREF=gi:11056051 /UG=Hs.78864 Fc fragment of IgG, low affinity IIa, receptor for (CD32) /FL=gb:NM_021642.1 gb:M31932.1 gb:J03619.1 gb:M28697.1	NM_021642		NP_067674

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
203566_s_at	0.005278	gb:NM_000645.1 /DEF=Homo sapiens amylo-1,6-glucosidase, 4-alpha-glucanotransferase (glycogen debranching enzyme, glycogen storage disease type III) (AGL), transcript variant 5, mRNA. /FEA=mRNA /GEN=AGL /PROD=amylo-1,6-glucosidase,4-alpha-glucanotransferase isoform 2 /DB_XREF=gi:4557282 /UG=Hs.904 amylo-1,6-glucosidase, 4-alpha-glucanotransferase (glycogen debranching enzyme, glycogen storage disease type III) /FL=gb:M85168.1 gb:NM_000645.1	NM_000645		NP_000637
203573_s_at	0.030947	gb:NM_004581.1 /DEF=Homo sapiens Rab geranylgeranyltransferase, alpha subunit (RABGGTA), mRNA. /FEA=mRNA /GEN=RABGGTA /PROD=Rab geranylgeranyltransferase, alpha subunit /DB_XREF=gi:4759015 /UG=Hs.78920 Rab geranylgeranyltransferase, alpha subunit /FL=gb:NM_004581.1	NM_004581		NP_004572
203574_at	0.002446	gb:NM_005384.1 /DEF=Homo sapiens nuclear factor, interleukin 3 regulated (NFIL3), mRNA. /FEA=mRNA /GEN=NFIL3 /PROD=nuclear factor, interleukin 3 regulated /DB_XREF=gi:4885516 /UG=Hs.79334 nuclear factor, interleukin 3 regulated /FL=gb:U26173.1 gb:NM_005384.1	NM_005384		NP_005375

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
203583_at	0.001835	gb:NM_014044.1 /DEF=Homo sapiens DKFZP564G0222 protein (DKFZP564G0222), mRNA. /FEA=mRNA /GEN=DKFZP564G0222 /PROD=DKFZP564G0222 protein /DB_XREF=gi:7661609 /UG=Hs.13370 DKFZP564G0222 protein /FL=gb:AF077038.1 gb:AL080115.1 gb:NM_014044.1	NM_014044		NP_054763
203584_at	0.034006	gb:NM_014673.1 /DEF=Homo sapiens KIAA0103 gene product (KIAA0103), mRNA. /FEA=mRNA /GEN=KIAA0103 /PROD=KIAA0103 gene product /DB_XREF=gi:7661909 /UG=Hs.154387 KIAA0103 gene product /FL=gb:D14659.1 gb:NM_014673.1	NM_014673		NP_055488
203585_at	0.018128	gb:NM_007150.1 /DEF=Homo sapiens zinc finger protein 185 (LIM domain) (ZNF185), mRNA. /FEA=mRNA /GEN=ZNF185 /PROD=zinc finger protein 185 (LIM domain) /DB_XREF=gi:6005971 /UG=Hs.16622 zinc finger protein 185 (LIM domain) /FL=gb:NM_007150.1	NM_007150		NP_009081
203592_s_at	0.045918	gb:NM_005860.1 /DEF=Homo sapiens follistatin-like 3 (secreted glycoprotein) (FSTL3), mRNA. /FEA=mRNA /GEN=FSTL3 /PROD=follistatin-like 3 glycoprotein /DB_XREF=gi:5031700 /UG=Hs.25348 follistatin-like 3 (secreted glycoprotein) /FL=gb:U76702.1 gb:NM_005860.1	NM_005860		NP_005851

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203594_at	0.003842	gb:NM_003729.1 /DEF=Homo sapiens RNA 3-terminal phosphate cyclase (RPC), mRNA. /FEA=mRNA /GEN=RPC /PROD=RNA 3-terminal phosphate cyclase. /DB_XREF=gi:4506588 /UG=Hs.27076 RNA 3-terminal phosphate cyclase /FL=gb:NM_003729.1	NM_003729		NP_003720
203598_s_at	7.85E-04	Consensus includes gb:AK000979.1 /DEF=Homo sapiens cDNA FLJ10117 fis, clone HEMBA1002810, highly similar to Homo sapiens formin binding protein 21 mRNA. /FEA=mRNA /DB_XREF=gi:7021977 /UG=Hs.28307 WW domain binding protein 4 (formin binding protein 21) /FL=gb:AF071185.1 gb:NM_007187.2	NM_007187		NP_009118
203604_at	0.003305	KIAA0222 gene product	N38750	Hs.48450	NP_055458
203605_at	0.001852	gb:NM_003136.1 /DEF=Homo sapiens signal recognition particle 54kD (SRP54), mRNA. /FEA=mRNA /GEN=SRP54 /PROD=signal recognition particle 54kD. /DB_XREF=gi:4507214 /UG=Hs.49346 signal recognition particle 54kD /FL=gb:U51920.1 gb:BC000652.1 gb:BC003389.1 gb:NM_003136.1	NM_003136		NP_003127

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203613_s_at	6.66E-04	gb:NM_002493.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6 (17kD, B17) (NDUFB6), mRNA. /FEA=mRNA /GEN=NDUFB6 /PROD=NADH dehydrogenase (ubiquinone) 1 betasubcomplex, 6 (17kD, B17) /DB_XREF=gi:4505364 /UG=Hs.109646 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6 (17kD, B17) /FL=gb:AF035840.1 gb:NM_002493.1 gb:AF067167.1	NM_002493		NP_002484
203620_s_at	0.002724	gb:NM_014824.1 /DEF=Homo sapiens KIAA0769 gene product (KIAA0769), mRNA. /FEA=mRNA /GEN=KIAA0769 /PROD=KIAA0769 gene product /DB_XREF=gi:7662295 /UG=Hs.19056 KIAA0769 gene product /FL=gb:AB018312.1 gb:NM_014824.1	NM_014824		NP_055639
203630_s_at	0.004987	gb:NM_006348.1 /DEF=Homo sapiens golgi transport complex 1 (90 kDa subunit) (GOLTC1), mRNA. /FEA=mRNA /GEN=GOLTC1 /PROD=golgi transport complex 1 (90 kDa subunit) /DB_XREF=gi:5453669 /UG=Hs.239631 golgi transport complex 1 (90 kDa subunit) /FL=gb:AF058718.1 gb:NM_006348.1	NM_006348		NP_006339
203633_at	0.025234	carnitine palmitoyltransferase 1A (liver)	BF001714	Hs.408050	NP_001867

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203635_at	7.25E-04	gb:NM_006052.1 /DEF=Homo sapiens Down syndrome critical region gene 3 (DSCR3), mRNA. /FEA=mRNA /GEN=DSCR3 /PROD=Down syndrome critical region protein 3 /DB_XREF=gi:5174424 /UG=Hs.26146 Down syndrome critical region gene 3 /FL=gb:D87343.1 gb:NM_006052.1	NM_006052		NP_006043
203643_at	0.016796	gb:NM_006494.1 /DEF=Homo sapiens Ets2 repressor factor (ERF), mRNA. /FEA=mRNA /GEN=ERF /PROD=Ets2 repressor factor /DB_XREF=gi:5729813 /UG=Hs.333069 Ets2 repressor factor /FL=gb:U15655.1 gb:NM_006494.1	NM_006494		NP_006485
203645_s_at	0.008944	gb:NM_004244.1 /DEF=Homo sapiens CD163 antigen (CD163), mRNA. /FEA=mRNA /GEN=CD163 /PROD=CD163 antigen /DB_XREF=gi:4758721 /UG=Hs.74076 CD163 antigen /FL=gb:NM_004244.1	NM_004244		NP_004235
203655_at	0.001719	gb:NM_006297.1 /DEF=Homo sapiens X-ray repair complementing defective repair in Chinese hamster cells 1 (XRCC1), mRNA. /FEA=mRNA /GEN=XRCC1 /PROD=X-ray repair cross complementing protein 1 /DB_XREF=gi:5454171 /UG=Hs.98493 X-ray repair complementing defective repair in Chinese hamster cells 1 /FL=gb:M36089.1 gb:NM_006297.1	NM_006297		NP_006288

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203656_at	0.001406	gb:NM_014845.1 /DEF=Homo sapiens KIAA0274 gene product (KIAA0274), mRNA. /FEA=mRNA /GEN=KIAA0274 /PROD=KIAA0274 gene product /DB_XREF=gi:7662033 /UG=Hs.10037 KIAA0274 gene product /FL=gb:D87464.1 gb:NM_014845.1	NM_014845		NP_055660
203658_at	0.010713	gb:BC001689.1 /DEF=Homo sapiens, carnitineacylcarnitine translocase, clone MGC:1207, mRNA, complete cds. /FEA=mRNA. /PROD=carnitineacylcarnitine translocase /DB_XREF=gi:12804552 /UG=Hs.13845 solute carrier family 25 (carnitineacylcarnitine translocase), member 20 /FL=gb:BC001689.1 gb:NM_000387.2	BC001689		NP_000378
203661_s_at	0.013163	gb:BC002660.1 /DEF=Homo sapiens, tropomodulin, clone MGC:3643, mRNA, complete cds. /FEA=mRNA /PROD=tropomodulin /DB_XREF=gi:12803650 /UG=Hs.170453 tropomodulin /FL=gb:BC002660.1 gb:M77016.1 gb:NM_003275.1	BC002660		NP_003266
203663_s_at	0.02217	gb:NM_004255.1 /DEF=Homo sapiens cytochrome c oxidase subunit Va (COX5A), mRNA. /FEA=mRNA /GEN=COX5A /PROD=cytochrome c oxidase subunit Va /DB_XREF=gi:4758037 /UG=Hs.181028 cytochrome c oxidase subunit Va /FL=gb:NM_004255.1 gb:M22760.1	NM_004255		NP_004246

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203665_at	0.005347	gb:NM_002133.1 /DEF=Homo sapiens heme oxygenase (decycling) 1 (HMOX1), mRNA. /FEA=mRNA /GEN=HMOX1 /PROD=heme oxygenase (decycling) 1 /DB_XREF=gi:4504436 /UG=Hs.202833 heme oxygenase (decycling) 1 /FL=gb:NM_002133.1	NM_002133		NP_002124
203667_at	0.031624	gb:NM_004607.1 /DEF=Homo sapiens tubulin-specific chaperone a (TBCA), mRNA. /FEA=mRNA /GEN=TBCA /PROD=beta-tubulin cofactor A /DB_XREF=gi:4759211 /UG=Hs.24930 tubulin-specific chaperone a /FL=gb:AF038952.1 gb:NM_004607.1	NM_004607		NP_004598
203668_at	0.022601	gb:NM_006715.1 /DEF=Homo sapiens mannosidase, alpha, class 2C, member 1 (MAN2C1), mRNA. /FEA=mRNA /GEN=MAN2C1 /PROD=mannosidase, alpha 6A8 /DB_XREF=gi:6631092 /UG=Hs.26232 mannosidase, alpha, class 2C, member 1 /FL=gb:U37248.1 gb:AF044414.2 gb:NM_006715.1	NM_006715		NP_006706
203669_s_at	0.025377	gb:NM_012079.2 /DEF=Homo sapiens diacylglycerol O-acyltransferase (mouse) homolog (DGAT), mRNA. /FEA=mRNA /GEN=DGAT /PROD=diacylglycerol O-acyltransferase (mouse)homolog /DB_XREF=gi:7382489 /UG=Hs.288627 diacylglycerol O-acyltransferase (mouse) homolog /FL=gb:AF059202.1 gb:NM_012079.2	NM_012079		NP_036211

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203672_x_at	0.01812	gb:U12387.1 /DEF=Human thiopurine methyltransferase (TPMT) mRNA, complete cds. /FEA=mRNA /GEN=TPMT /PROD=thiopurine methyltransferase /DB_XREF=gi:805083 /UG=Hs.296922 thiopurine S-methyltransferase /FL=gb:BC005339.1 gb:NM_000367.1 gb:U12387.1	U12387		NP_000358
203675_at	6.66E-04	gb:NM_005013.1 /DEF=Homo sapiens nucleobindin 2 (NUCB2), mRNA. /FEA=mRNA /GEN=NUCB2 /PROD=nucleobindin 2 /DB_XREF=gi:4826869 /UG=Hs.3164 nucleobindin 2 /FL=gb:AF052642.1 gb:AF052643.1 gb:AF052644.1 gb:NM_005013.1	NM_005013		NP_005004
203686_at	0.027336	gb:NM_002434.1 /DEF=Homo sapiens N-methylpurine-DNA glycosylase (MPG), mRNA. /FEA=mRNA /GEN=MPG /PROD=N-methylpurine-DNA glycosylase /DB_XREF=gi:4505232 /UG=Hs.79396 N-methylpurine-DNA glycosylase /FL=gb:M74905.1 gb:L10752.1 gb:NM_002434.1	NM_002434		NP_002425
203690_at	0.001179	gb:NM_006322.1 /DEF=Homo sapiens spindle pole body protein (GCP3), mRNA. /FEA=mRNA /GEN=GCP3 /PROD=spindle pole body protein /DB_XREF=gi:5453659 /UG=Hs.9884 spindle pole body protein /FL=gb:AF042378.1 gb:NM_006322.1	NM_006322		NP_006313

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
203691_at	0.011969	gb:NM_002638.1 /DEF=Homo sapiens protease inhibitor 3, skin-derived (SKALP) (PI3), mRNA. /FEA=mRNA /GEN=PI3 /PROD=protease inhibitor 3, skin-derived (SKALP). /DB_XREF=gi:4505786 /UG=Hs.112341 protease inhibitor 3, skin-derived (SKALP) /FL=gb:NM_002638.1	NM_002638		NP_002629
203693_s_at	0.008506	gb:NM_001949.2 /DEF=Homo sapiens E2F transcription factor 3 (E2F3) mRNA, complete cds. /FEA=mRNA /GEN=E2F3 /PROD=E2F transcription factor 3 /DB_XREF=gi:12669913 /UG=Hs.1189 E2F transcription factor 3 /FL=gb:NM_001949.2	NM_001949		NP_001940
203704_s_at	0.001763	ras responsive element binding protein 1	AW118862	Hs.171942	NP_002946
203707_at	0.032443	gb:NM_005741.1 /DEF=Homo sapiens zinc finger protein 263 (ZNF263), mRNA. /FEA=mRNA /GEN=ZNF263 /PROD=zinc finger protein 263 /DB_XREF=gi:5032240 /UG=Hs.182528 zinc finger protein 263 /FL=gb:D88827.1 gb:NM_005741.1	NM_005741		NP_005732
203713_s_at	0.003584	gb:NM_004524.1 /DEF=Homo sapiens lethal giant larvae (Drosophila) homolog 2 (LLGL2), mRNA. /FEA=mRNA /GEN=LLGL2 /PROD=lethal giant larvae (Drosophila) homolog 2 /DB_XREF=gi:4758679 /UG=Hs.3123 lethal giant larvae (Drosophila) homolog 2 /FL=gb:NM_004524.1	NM_004524		NP_004515

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203721_s_at	0.01063	gb:NM_016001.1 /DEF=Homo sapiens CGI-48 protein (LOC51096), mRNA. /FEA=mRNA /GEN=LOC51096 /PROD=CGI-48 protein /DB_XREF=gi:7705764 /UG=Hs.6153 CGI-48 protein /FL=gb:AF151806.1 gb:NM_016001.1	NM_016001		NP_057085
203725_at	7.63E-04	gb:NM_001924.2 /DEF=Homo sapiens growth arrest and DNA-damage-inducible, alpha (GADD45A), mRNA. /FEA=mRNA /GEN=GADD45A /PROD=growth arrest and DNA-damage-inducible, alpha /DB_XREF=gi:9790904 /UG=Hs.80409 growth arrest and DNA-damage-inducible, alpha /FL=gb:M60974.1 gb:NM_001924.2	NM_001924		NP_001915
203731_s_at	0.019689	gb:NM_014569.1 /DEF=Homo sapiens zinc finger protein homologous to Zfp95 in mouse (ZFP95), mRNA. /FEA=mRNA /GEN=ZFP95 /PROD=zinc finger protein homologous to Zfp95 in mouse /DB_XREF=gi:11036641 /UG=Hs.110839 zinc finger protein homologous to Zfp95 in mouse /FL=gb:NM_014569.1 gb:AB023232.1	NM_014569		NP_659570
203732_at	0.005347	gb:NM_016213.1 /DEF=Homo sapiens thyroid hormone receptor interactor 4 (TRIP4), mRNA. /FEA=mRNA /GEN=TRIP4 /PROD=activating signal cointegrator 1 /DB_XREF=gi:7706430 /UG=Hs.116784 thyroid hormone receptor interactor 4 /FL=gb:AF168418.1 gb:NM_016213.1	NM_016213		NP_057297

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203739_at	0.00121	gb:NM_006526.1 /DEF=Homo sapiens zinc finger protein 217 (ZNF217), mRNA. /FEA=mRNA /GEN=ZNF217 /PROD=zinc finger protein 217 /DB_XREF=gi:5730123 /UG=Hs.155040 zinc finger protein 217 /FL=gb:AF041259.1 gb:NM_006526.1	NM_006526		NP_006517
203743_s_at	0.027672	gb:NM_003211.1 /DEF=Homo sapiens thymine-DNA glycosylase (TDG), mRNA. /FEA=mRNA /GEN=TDG /PROD=thymine-DNA glycosylase /DB_XREF=gi:4507422 /UG=Hs.173824 thymine-DNA glycosylase /FL=gb:U51166.1 gb:NM_003211.1	NM_003211		NP_003202
203744_at	0.001916	gb:NM_005342.1 /DEF=Homo sapiens high-mobility group (nonhistone chromosomal) protein 4 (HMG4), mRNA. /FEA=mRNA /GEN=HMG4 /PROD=high-mobility group (nonhistone chromosomal) protein 4 /DB_XREF=gi:4885420 /UG=Hs.19114 high-mobility group (nonhistone chromosomal) protein 4 /FL=gb:NM_005342.1	NM_005342		NP_005333
203746_s_at	0.007022	gb:NM_005333.1 /DEF=Homo sapiens holocytochrome c synthase (cytochrome c heme-lyase) (HCCS), mRNA. /FEA=mRNA /GEN=HCCS /PROD=holocytochrome c synthase (cytochrome heme-lyase) /DB_XREF=gi:4885400 /UG=Hs.211571 holocytochrome c synthase (cytochrome c heme-lyase) /FL=gb:U36787.1 gb:NM_005333.1	NM_005333		NP_005324

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203748_x_at	9.85E-04	gb:NM_016839.1 /DEF=Homo sapiens RNA binding motif, single stranded interacting protein 1 (RBMS1), transcript variant M5SP-2, mRNA. /FEA=mRNA /GEN=RBMS1 /PROD=RNA binding motif, single stranded interacting protein 1, isoform b /DB_XREF=gi:8400723 /UG=Hs.241567 RNA binding motif, single stranded interacting protein 1 /FL=gb:NM_016839.1	NM_016839		NP_058523
203757_s_at	0.025095	gb:BC005008.1 /DEF=Homo sapiens, carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen), clone MGC:10467, mRNA, complete cds. /FEA=mRNA /PROD=carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen) /DB_XREF=gi:13477106 /UG=Hs.73848 carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen) /FL=gb:BC005008.1 gb:M18216.1 gb:M29541.1 gb:NM_002483.1	BC005008		NP_002474
203761_at	0.00643	gb:NM_006748.1 /DEF=Homo sapiens Src-like-adapter (SLA), mRNA. /FEA=mRNA /GEN=SLA /PROD=Src-like-adapter /DB_XREF=gi:5803170 /UG=Hs.75367 Src-like-adapter /FL=gb:U30473.1 gb:D89077.1 gb:U44403.1 gb:NM_006748.1	NM_006748		NP_006739

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
203764_at	0.013163	gb:NM_014750.1 /DEF=Homo sapiens KIAA0008 gene product (KIAA0008), mRNA. /FEA=mRNA /GEN=KIAA0008 /PROD=KIAA0008 gene product /DB_XREF=gi:7661851 /UG=Hs.77695 KIAA0008 gene product /FL=gb:D13633.1 gb:NM_014750.1	NM_014750		NP_055565
203765_at	0.032771	gb:NM_012198.1 /DEF=Homo sapiens grancalcin (GCL), mRNA. /FEA=mRNA /GEN=GCL /PROD=grancalcin /DB_XREF=gi:6912387 /UG=Hs.79381 grancalcin /FL=gb:BC005214.1 gb:M81637.1 gb:NM_012198.1	NM_012198		NP_036330
203773_x_at	0.048412	gb:NM_000712.1 /DEF=Homo sapiens biliverdin reductase A (BLVRA), mRNA. /FEA=mRNA /GEN=BLVRA /PROD=biliverdin reductase A /DB_XREF=gi:4502416 /UG=Hs.81029 biliverdin reductase A /FL=gb:U34877.1 gb:NM_000712.1	NM_000712		NP_000703
203775_at	0.001763	gb:NM_014251.1 /DEF=Homo sapiens solute carrier family 25, member 13 (citrin) (SLC25A13), mRNA. /FEA=mRNA /GEN=SLC25A13 /PROD=solute carrier family 25, member 13 (citric) /DB_XREF=gi:7657580 /UG=Hs.9599 solute carrier family 25, member 13 (citric) /FL=gb:AF118838.1 gb:NM_014251.1	NM_014251		NP_055066

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203778_at	0.002559	gb:NM_005908.1 /DEF=Homo sapiens mannosidase, beta A, lysosomal (MANBA), mRNA. /FEA=mRNA /GEN=MANBA /PROD=mannosidase, beta A, lysosomal /DB_XREF=gi:5174522 /UG=Hs.115945 mannosidase, beta A, lysosomal /FL=gb:U60337.1 gb:NM_005908.1	NM_005908		NP_005899
203781_at	0.014263	gb:NM_004891.1 /DEF=Homo sapiens chromosome 2 open reading frame 1 (C2ORF1), mRNA. /FEA=mRNA /GEN=C2ORF1 /PROD=chromosome 2 open reading frame 1 /DB_XREF=gi:4759047 /UG=Hs.14454 chromosome 2 open reading frame 1 /FL=gb:AF047440.1 gb:NM_004891.1	NM_004891		NP_663303
203799_at	0.018248	gb:NM_014880.1 /DEF=Homo sapiens KIAA0022 gene product (KIAA0022), mRNA. /FEA=mRNA /GEN=KIAA0022 /PROD=KIAA0022 gene product /DB_XREF=gi:7661867 /UG=Hs.2441 KIAA0022 gene product /FL=gb:D14664.1 gb:NM_014880.1	NM_014880		NP_055695
203800_s_at	0.002749	Consensus includes gb:BG254653 /FEA=EST /DB_XREF=gi:12764469 /DB_XREF=est:602368621F1 /CLONE=IMAGE:4476773 /UG=Hs.247324 hypothetical protein LOC63931 /FL=gb:NM_022100.1	NM_022100		NP_071383

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203814_s_at	0.007652	gb:NM_000904.1 /DEF=Homo sapiens NAD(P)H menadione oxidoreductase 2, dioxin-inducible (NMOR2), mRNA. /FEA=mRNA /GEN=NMOR2 /PROD=NAD(P)H menadione oxidoreductase 2,dioxin-inducible /DB_XREF=gi:4505416 /UG=Hs.73956 NAD(P)H menadione oxidoreductase 2, dioxin-inducible /FL=gb:J02888.1 gb:NM_000904.1	NM_000904		NP_000895
203825_at	0.01132	gb:NM_007371.2 /DEF=Homo sapiens bromodomain-containing 3 (BRD3), mRNA. /FEA=mRNA /GEN=BRD3 /PROD=bromodomain-containing protein 3 /DB_XREF=gi:12408642 /UG=Hs.86896 bromodomain-containing 3 /FL=gb:NM_007371.2 gb:D26362.1	NM_007371		NP_031397
203827_at	0.00236	gb:NM_017983.1 /DEF=Homo sapiens hypothetical protein FLJ10055 (FLJ10055), mRNA. /FEA=mRNA /GEN=FLJ10055 /PROD=hypothetical protein FLJ10055 /DB_XREF=gi:8922207 /UG=Hs.9398 hypothetical protein FLJ10055 /FL=gb:NM_017983.1	NM_017983		NP_060453
203837_at	0.014263	gb:NM_005923.2 /DEF=Homo sapiens mitogen-activated protein kinase kinase kinase 5 (MAP3K5), mRNA. /FEA=mRNA /GEN=MAP3K5 /PROD=MAPERK kinase kinase 5 /DB_XREF=gi:6031181 /UG=Hs.151988 mitogen-activated protein kinase kinase kinase 5 /FL=gb:U67156.1 gb:D84476.1 gb:NM_005923.2	NM_005923		NP_005914

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203856_at	0.038618	gb:NM_003384.1 /DEF=Homo sapiens vaccinia related kinase 1 (VRK1), mRNA. /FEA=mRNA /GEN=VRK1 /PROD=vaccinia related kinase 1 /DB_XREF=gi:4507902 /UG=Hs.48269 vaccinia related kinase 1 /FL=gb:AB000449.1 gb:NM_003384.1	NM_003384		NP_003375
203857_s_at	0.006157	gb:NM_006810.1 /DEF=Homo sapiens for protein disulfide isomerase-related (PDIR), mRNA. /FEA=mRNA /GEN=PDIR /PROD=for protein disulfide isomerase-related /DB_XREF=gi:5803120 /UG=Hs.76901 for protein disulfide isomerase-related /FL=gb:D49490.1 gb:NM_006810.1	NM_006810		NP_006801
203880_at	0.004987	gb:NM_005694.1 /DEF=Homo sapiens COX17 (yeast) homolog, cytochrome c oxidase assembly protein (COX17), mRNA. /FEA=mRNA /GEN=COX17 /PROD=COX17 (yeast) homolog, cytochrome c oxidase assembly protein /DB_XREF=gi:5031644 /UG=Hs.16297 COX17 (yeast) homolog, cytochrome c oxidase assembly protein /FL=gb:L77701.1 gb:NM_005694.1	NM_005694		NP_005685
203884_s_at	0.006043	gb:NM_014904.1 /DEF=Homo sapiens KIAA0941 protein (KIAA0941), mRNA. /FEA=mRNA /GEN=KIAA0941 /PROD=KIAA0941 protein /DB_XREF=gi:7662393 /UG=Hs.173656 KIAA0941 protein /FL=gb:AB023158.1 gb:NM_014904.1	NM_014904		NP_055719

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
203885_at	7.25E-04	gb:NM_014999.1 /DEF=Homo sapiens KIAA0118 protein (KIAA0118); mRNA. /FEA=mRNA /GEN=KIAA0118 /PROD=KIAA0118 protein /DB_XREF=gi:7661921. /UG=Hs.184627 KIAA0118 protein /FL=gb:AF091035.1 gb:NM_014999.1	NM_014999		NP_055814
203887_s_at	0.003373	gb:NM_000361.1 /DEF=Homo sapiens thrombomodulin (THBD), mRNA. /FEA=mRNA /GEN=THBD /PROD=thrombomodulin /DB_XREF=gi:4507482. /UG=Hs.2030 thrombomodulin /FL=gb:M16552.1 gb:NM_000361.1	NM_000361		NP_000352
203888_at	0.003248	gb:NM_000361.1 /DEF=Homo sapiens thrombomodulin (THBD), mRNA. /FEA=mRNA /GEN=THBD /PROD=thrombomodulin /DB_XREF=gi:4507482. /UG=Hs.2030 thrombomodulin /FL=gb:M16552.1 gb:NM_000361.1	NM_000361		NP_000352
203897_at	0.003009	hypothetical protein A-211C6.1	BE963444	Hs.28607	NP_065157
203904_x_at	0.017522	gb:NM_002231.2. /DEF=Homo sapiens kangai 1 (suppression of tumorigenicity 6; prostate; CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4)) (KAI1), mRNA. /FEA=mRNA /GEN=KAI1 /PROD=kangai 1 /DB_XREF=gi:13259537. /UG=Hs.323949 kangai 1 (suppression of tumorigenicity 6; prostate; CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4)) /FL=gb:BC000726.1 gb:BC001821.1 gb:NM_002231.2 gb:U20770.1	NM_002231		NP_002222

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203906_at	7.63E-04	KIAA0763 gene product	AI652645	Hs.409124	NP_055684
203907_s_at	7.63E-04	gb:NM_014869.1 /DEF=Homo sapiens KIAA0763 gene product (KIAA0763), mRNA. /FEA=mRNA /GEN=KIAA0763 /PROD=KIAA0763 gene product /DB_XREF=gi:7662289 /UG=Hs.4764 KIAA0763 gene product /FL=gb:AB018306.1 gb:NM_014869.1	NM_014869		NP_055684
203909_at	7.63E-04	gb:NM_006359.1 /DEF=Homo sapiens solute carrier family 9 (sodiumhydrogen exchanger), isoform 6 (SLC9A6), mRNA. /FEA=mRNA /GEN=SLC9A6 /PROD=solute carrier family 9 (sodiumhydrogenexchanger), isoform 6 /DB_XREF=gi:5454069 /UG=Hs.62185 solute carrier family 9 (sodiumhydrogen exchanger), isoform 6 /FL=gb:AF030409.1 gb:NM_006359.1	NM_006359		NP_006350
203912_s_at	7.83E-04	gb:NM_006730.1 /DEF=Homo sapiens deoxyribonuclease I-like 1 (DNASE1L1), mRNA. /FEA=mRNA /GEN=DNASE1L1 /PROD=deoxyribonuclease I-like 1 /DB_XREF=gi:5803006 /UG=Hs.77091 deoxyribonuclease I-like 1 /FL=gb:U06846.1 gb:NM_006730.1	NM_006730		NP_006721

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203921_at	0.019976	gb:NM_004267.1 /DEF=Homo sapiens carbohydrate (chondroitin 6keratan) sulfotransferase 2 (CHST2), mRNA. /FEA=mRNA /GEN=CHST2 /PROD=carbohydrate (chondroitin 6keratan)sulfotransferase 2 /DB_XREF=gi:4757983 /UG=Hs.8786 carbohydrate (chondroitin 6keratan) sulfotransferase 2 /FL=gb:AB021124.1 gb:AB014679.1 gb:AB014680.1 gb:AF083066.1 gb:NM_004267.1	NM_004267		NP_004258
203925_at	7.63E-04	gb:NM_002061.1 /DEF=Homo sapiens glutamate-cysteine ligase, modifier subunit (GCLM), mRNA. /FEA=mRNA /GEN=GCLM /PROD=glutamate-cysteine ligase regulatory protein /DB_XREF=gi:4504010 /UG=Hs.89709 glutamate-cysteine ligase, modifier subunit /FL=gb:NM_002061.1 gb:L35546.1	NM_002061		NP_002052
203936_s_at	0.009678	gb:NM_004994.1 /DEF=Homo sapiens matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase) (MMP9), mRNA. /FEA=mRNA /GEN=MMP9 /PROD=matrix metalloproteinase 9 preproprotein /DB_XREF=gi:4826835 /UG=Hs.151738 matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase) /FL=gb:J05070.1 gb:NM_004994.1	NM_004994		NP_004985

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
203945_at	0.004648	gb:NM_001172.2 /DEF=Homo sapiens arginase, type II (ARG2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=ARG2 /PROD=arginase, type II precursor /DB_XREF=gi:10947110 /UG=Hs.172851 arginase, type II /FL=gb:NM_001172.2 gb:BC001350.1 gb:D86724.1 gb:U75667.1 gb:U82256.1	NM_001172		NP_001163
203958_s_at	0.028228	KIAA0478 gene product	AI557467	Hs.4236	NP_055685
203960_s_at	0.024046	gb:NM_016126.1 /DEF=Homo sapiens HSPCO34 protein (LOC51668), mRNA. /FEA=mRNA /GEN=LOC51668 /PROD=HSPCO34 protein /DB_XREF=gi:7706382 /UG=Hs.46967 HSPCO34 protein /FL=gb:BC005245.1 gb:AF100747.1 gb:NM_016126.1	NM_016126		NP_057210
203966_s_at	6.66E-04	gb:NM_021003.1 /DEF=Homo sapiens protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform (PPM1A), mRNA. /FEA=mRNA /GEN=PPM1A /PROD=protein phosphatase 1A (formerly 2C),magnesium-dependent, alpha isoform /DB_XREF=gi:10337594 /UG=Hs.57764 protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform /FL=gb:NM_021003.1	NM_021003		NP_808821

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203971_at	0.038618	gb:NM_001859.1 /DEF=Homo sapiens solute carrier family 31 (copper transporters), member 1 (SLC31A1), mRNA. /FEA=mRNA /GEN=SLC31A1. /PROD=solute carrier family 31 (copper transporters), member 1 /DB_XREF=gi:4507014 /UG=Hs.73614 solute carrier family 31 (copper transporters), member 1 /FL=gb:U83460.1 gb:NM_001859.1	NM_001859		NP_001850
203974_at	0.01289	gb:NM_012080.1 /DEF=Homo sapiens DNA segment, numerous copies, expressed probes (GS1 gene) (DXF68S1E), mRNA. /FEA=mRNA /GEN=DXF68S1E. /PROD=DNA segment, numerous copies, expressed probes(GS1 gene) /DB_XREF=gi:6912345 /UG=Hs.78991 DNA segment, numerous copies, expressed probes (GS1 gene) /FL=gb:M86934.1 gb:NM_012080.1	NM_012080		NP_036212
203983_at	0.01412	gb:NM_005999.1 /DEF=Homo sapiens translin-associated factor X (TSNAX), mRNA. /FEA=mRNA /GEN=TSNAX /PROD=translin-associated factor X /DB_XREF=gi:5174730 /UG=Hs.96247 translin-associated factor X /FL=gb:NM_005999.1	NM_005999		NP_005990

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Prot in Accession No.
203984_s_at	0.020419	gb:U60521.1 /DEF=Human protease proMch6 (Mch6) mRNA, complete cds. /FEA=mRNA /GEN=Mch6 /PROD=proMch6 /DB_XREF=gi:1532150 /UG=Hs.100641 caspase 9, apoptosis-related cysteine protease /FL=gb:BC002452.1 gb:U56390.1 gb:U60521.1 gb:NM_001229.1	U60521		NP_127463
203985_at	0.028971	gb:NM_012256.1 /DEF=Homo sapiens zinc finger protein 212 (ZNF212), mRNA. /FEA=mRNA /GEN=ZNF212 /PROD=zinc finger protein 212 /DB_XREF=gi:6912749 /UG=Hs.108139 zinc finger protein 212 /FL=gb:U38864.1 gb:NM_012256.1	NM_012256		NP_036388
203992_s_at	0.006722	gb:AF000992.1 /DEF=Homo sapiens ubiquitous TPR motif, X isoform (UTX) mRNA, alternative transcript 1, complete cds. /FEA=mRNA /GEN=UTX /PROD=ubiquitous TPR motif, X isoform /DB_XREF=gi:2580569 /UG=Hs.13980 ubiquitously transcribed tetratricopeptide repeat gene, X chromosome /FL=gb:NM_021140.1 gb:AF000992.1 gb:AF000993.1	AF000992		NP_066963

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniGene Accession No.	Protein Accession No.
204001_at	0.036689	gb:NM_003084.1 /DEF=Homo sapiens small nuclear RNA activating complex, polypeptide 3, 50kD (SNAPC3), mRNA. /FEA=mRNA /GEN=SNAPC3 /PROD=small nuclear RNA activating complex, polypeptide 3, 50kD /DB_XREF=gi:4507104 /UG=Hs.164915 small nuclear RNA activating complex, polypeptide 3, 50kD /FL=gb:U66413.1 gb:NM_003084.1	NM_003084		NP_003075
204009_s_at	7.25E-04	v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog	W80678	Hs.351221	NP_203524
204018_x_at	0.001942	gb:NM_000558.2 /DEF=Homo sapiens hemoglobin, alpha 1 (HBA1), mRNA. /FEA=mRNA /GEN=HBA1 /PROD=hemoglobin, alpha 1. /DB_XREF=gi:6715603 /UG=Hs.251577 hemoglobin, alpha 1 /FL=gb:NM_000558.2	NM_000558		NP_000549
204022_at	6.66E-04	Nedd-4-like ubiquitin-protein ligase	AI668780	Hs.333382	NP_008945
204026_s_at	0.00121	gb:NM_007057.1 /DEF=Homo sapiens ZW10 interactor (ZWINT), mRNA. /FEA=mRNA /GEN=ZWINT /PROD=ZW10 interactor /DB_XREF=gi:6857828 /UG=Hs.42650 ZW10 interactor /FL=gb:AF067656.1 gb:NM_007057.1	NM_007057		NP_127490

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204028_s_at	0.001406	gb:NM_012197.2 /DEF=Homo sapiens rab6 GTPase activating protein (GAP and centrosome-associated) (GAPCENA), mRNA. /FEA=mRNA /GEN=GAPCENA /PROD=rab6 GTPase activating protein (GAP and centrosome-associated) /DB_XREF=gi:12232372 /UG=Hs.55099 rab6 GTPase activating protein (GAP and centrosome-associated) /FL=gb:NM_012197.2	NM_012197		NP_036329
204031_s_at	0.008874	gb:NM_005016.1 /DEF=Homo sapiens poly(rC)-binding protein 2 (PCBP2), mRNA. /FEA=mRNA /GEN=PCBP2 /PROD=poly(rC)-binding protein 2 /DB_XREF=gi:4826885 /UG=Hs.63525 poly(rC)-binding protein 2 /FL=gb:BC001155.1 gb:NM_005016.1	NM_005016		NP_114366
204034_at	6.66E-04	gb:NM_014297.1 /DEF=Homo sapiens protein expressed in thyroid (YF13H12), mRNA. /FEA=mRNA /GEN=YF13H12 /PROD=protein expressed in thyroid. /DB_XREF=gi:7657686 /UG=Hs.7486 protein expressed in thyroid /FL=gb:NM_014297.1	NM_014297		NP_055112

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204038_s_at	0.02217	gb:NM_001401.1 /DEF=Homo sapiens endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2 (EDG2), mRNA. /FEA=mRNA /GEN=EDG2 /PROD=endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2 /DB_XREF=gi:4503456 /UG=Hs.75794 endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2 /FL=gb:U78192.1 gb:U80811.1 gb:NM_001401.1	NM_001401		NP_476500
204050_s_at	7.76E-04	gb:NM_001833.1 /DEF=Homo sapiens clathrin, light polypeptide (Lca) (CLTA), transcript variant nonbrain, mRNA. /FEA=mRNA /GEN=CLTA /PROD=clathrin, light polypeptide A (Lca) isoform a /DB_XREF=gi:4502898 /UG=Hs.104143 clathrin, light polypeptide (Lca) /FL=gb:M20472.1 gb:NM_001833.1	NM_001833		NP_009027
204079_at	0.001406	gb:NM_003595.1 /DEF=Homo sapiens tyrosylprotein sulfotransferase 2 (TPST2), mRNA. /FEA=mRNA /GEN=TPST2 /PROD=tyrosylprotein sulfotransferase 2 /DB_XREF=gi:4507666 /UG=Hs.26350 tyrosylprotein sulfotransferase 2 /FL=gb:AL136623.1 gb:BC001057.1 gb:AF061254.1 gb:AF049891.1 gb:NM_003595.1	NM_003595		NP_003586

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204082_at	0.005347	gb:NM_006195.1 /DEF=Homo sapiens pre-B-cell leukemia transcription factor 3 (PBX3), mRNA. /FEA=mRNA /GEN=PBX3 /PROD=pre-B-cell leukemia transcription factor 3 /DB_XREF=gi:5453851 /UG=Hs.294101 pre-B-cell leukemia transcription factor 3 /FL=gb:NM_006195.1	NM_006195		NP_006186
204091_at	0.018701	gb:NM_002601.1 /DEF=Homo sapiens phosphodiesterase 6D, cGMP-specific, rod, delta (PDE6D), mRNA. /FEA=mRNA /GEN=PDE6D /PROD=phosphodiesterase 6D, cGMP-specific, rod, delta /DB_XREF=gi:4505670 /UG=Hs.48291 phosphodiesterase 6D, cGMP-specific, rod, delta /FL=gb:AF022912.1 gb:NM_002601.1	NM_002601		NP_002592
204093_at	0.036998	gb:NM_001239.1 /DEF=Homo sapiens cyclin H (CCNH), mRNA. /FEA=mRNA /GEN=CCNH /PROD=cyclin H /DB_XREF=gi:4502622 /UG=Hs.514 cyclin H /FL=gb:BC005280.1 gb:NM_001239.1 gb:U12685.1 gb:U11791.1	NM_001239		NP_001230

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	UniProt Accession No.	Protein Accession No.
204099_at	0.010277	gb:NM_003078.1 /DEF=Homo sapiens SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3 (SMARCD3), mRNA. /FEA=mRNA /GEN=SMARCD3 /PROD=SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3. /DB_XREF=gi:4507086 /UG=Hs.71622 SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3 /FL=gb:BC002628.1 gb:U66619.1 gb:NM_003078.1	NM_003078		NP_003069
204116_at	0.010713	gb:NM_000206.1 /DEF=Homo sapiens interleukin 2 receptor, gamma (severe combined immunodeficiency) (IL2RG), mRNA. /FEA=mRNA /GEN=IL2RG /PROD=interleukin 2 receptor, gamma chain, precursor /DB_XREF=gi:4557881 /UG=Hs.84 interleukin 2 receptor, gamma (severe combined immunodeficiency) /FL=gb:NM_000206.1	NM_000206		NP_000197
204118_at	0.029485	gb:NM_001778.1 /DEF=Homo sapiens CD48 antigen (B-cell membrane protein) (CD48), mRNA. /FEA=mRNA /GEN=CD48 /PROD=CD48 antigen (B-cell membrane protein) /DB_XREF=gi:4502674 /UG=Hs.901 CD48 antigen (B-cell membrane protein) /FL=gb:M59904.1 gb:M37766.1 gb:NM_001778.1	NM_001778		NP_001769

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204125_at	0.019342	gb:NM_016013.1 /DEF=Homo sapiens CGI-65 protein (LOC51103), mRNA. /FEA=mRNA /GEN=LOC51103 /PROD=CGI-65 protein /DB_XREF=gi:7705778 /UG=Hs.106529 CGI-65 protein /FL=gb:BC000780.1 gb:AF151823.1 gb:NM_016013.1	NM_016013		NP_057097
204137_at	0.013589	gb:NM_003272.1 /DEF=Homo sapiens transmembrane 7 superfamily member 1 (upregulated in kidney) (TM7SF1), mRNA. /FEA=mRNA /GEN=TM7SF1 /PROD=transmembrane 7 superfamily member 1 (upregulated in kidney) /DB_XREF=gi:4507544 /UG=Hs.15791 transmembrane 7 superfamily member 1 (upregulated in kidney) /FL=gb:AF027826.1 gb:NM_003272.1	NM_003272		NP_003263
204150_at	0.023971	gb:NM_015136.1 /DEF=Homo sapiens KIAA0246 protein (KIAA0246), mRNA. /FEA=mRNA /GEN=KIAA0246 /PROD=KIAA0246 protein /DB_XREF=gi:12225239 /UG=Hs.301989 KIAA0246 protein /FL=gb:NM_015136.1	NM_015136		NP_055951
204157_s_at	0.001336	gb:NM_025164.1 /DEF=Homo sapiens KIAA0999 protein (KIAA0999), mRNA. /FEA=mRNA /GEN=KIAA0999 /PROD=hypothetical protein FLJ12240 /DB_XREF=gi:13386467 /UG=Hs.4278 KIAA0999 protein /FL=gb:NM_025164.1	NM_025164		NP_079440

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
204159_at	0.012837	gb:NM_001262.1 /DEF=Homo sapiens cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4) (CDKN2C), mRNA. /FEA=mRNA /GEN=CDKN2C /PROD=cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4) /DB_XREF=gi:4502750 /UG=Hs.4854 cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4) /FL=gb:AF041248.1 gb:NM_001262.1	NM_001262		NP_523240
204160_s_at	0.002109	ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function)	AW194947	Hs.54037	NP_055751
204161_s_at	0.040553	gb:NM_014936.1 /DEF=Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function) (ENPP4), mRNA. /FEA=mRNA /GEN=ENPP4 /PROD=ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function) /DB_XREF=gi:7662357 /UG=Hs.54037 ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function) /FL=gb:AB020686.1 gb:NM_014936.1	NM_014936		NP_055751
204169_at	0.01534	gb:NM_000883.1 /DEF=Homo sapiens IMP (inosine monophosphate) dehydrogenase 1 (IMPDH1), mRNA. /FEA=mRNA /GEN=IMPDH1 /PROD=IMP (inosine monophosphate) dehydrogenase 1 /DB_XREF=gi:4504686 /UG=Hs.850 IMP (inosine monophosphate) dehydrogenase 1 /FL=gb:J05272.1 gb:NM_000883.1	NM_000883		NP_000874

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
204174_at	0.018534	gb:NM_001629.1 /DEF=Homo sapiens arachidonate 5-lipoxygenase-activating protein (ALOX5AP), mRNA. /FEA=mRNA /GEN=ALOX5AP /PROD=arachidonate 5-lipoxygenase-activating protein /DB_XREF=gi:4502058 /UG=Hs.100194 arachidonate 5-lipoxygenase-activating protein /FL=gb:NM_001629.1	NM_001629		NP_001620
204177_s_at	0.007818	gb:NM_014458.2 /DEF=Homo sapiens Kelch motif containing protein (AB026190), mRNA. /FEA=mRNA /GEN=AB026190 /PROD=Kelch motif containing protein /DB_XREF=gi:13677223 /UG=Hs.106290 Kelch motif containing protein /FL=gb:NM_014458.2 gb:AB026190.1	NM_014458		NP_055273
204180_s_at	0.031624	Consensus includes gb:AI745225 /FEA=EST /DB_XREF=gi:5113513 /DB_XREF=est:wg10d12.x1 /CLONE=IMAGE:2364695 /UG=Hs.127649 KIAA0414 protein /FL=gb:NM_014007.1	NM_014007		NP_054726
204185_x_at	6.66E-04	gb:NM_005038.1 /DEF=Homo sapiens peptidylprolyl isomerase D (cyclophilin D) (PPID), mRNA. /FEA=mRNA /GEN=PPID /PROD=peptidylprolyl isomerase D (cyclophilin D) /DB_XREF=gi:4826931 /UG=Hs.143482 peptidylprolyl isomerase D (cyclophilin D) /FL=gb:L11667.1 gb:NM_005038.1	NM_005038		NP_005029
204186_s_at	0.014372	peptidylprolyl isomerase D (cyclophilin D)	AI014573	Hs.143482	NP_005029
204198_s_at	0.002214	runt-related transcription factor 3	AA541630	Hs.170019	NP_004341

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gen Accession No.	Unigen Accession No.	Protein Accession No.
204204_at	0.007693	gb:NM_001860.1 /DEF=Homo sapiens solute carrier family 31 (copper transporters), member 2 (SLC31A2), mRNA. /FEA=mRNA /GEN=SLC31A2 /PROD=solute carrier family 31 (copper transporters),member 2 /DB_XREF=gi:4507016 /UG=Hs.24030 solute carrier family 31 (copper transporters), member 2 /FL=gb:U83461.1 gb:NM_001860.1	NM_001860		NP_001851
204206_at	0.007818	gb:NM_020310.1 /DEF=Homo sapiens MAX binding protein (MNT), mRNA. /FEA=mRNA /GEN=MNT /PROD=MAX binding protein /DB_XREF=gi:9945317 /UG=Hs.25497 MAX binding protein /FL=gb:NM_020310.1	NM_020310		NP_064706
204209_at	0.044364	phosphate cytidyltransferase 1, choline, alpha isoform	AI638771	Hs.273558	
204214_s_at	7.41E-04	gb:NM_006834.1 /DEF=Homo sapiens RAB32, member RAS oncogene family (RAB32), mRNA. /FEA=mRNA /GEN=RAB32 /PROD=RAB32, member RAS oncogene family /DB_XREF=gi:5803132 /UG=Hs.32217 RAB32, member RAS oncogene family /FL=gb:U71127.1 gb:NM_006834.1	NM_006834		NP_006825
204216_s_at	0.048867	gb:NM_024824.1 /DEF=Homo sapiens hypothetical protein FLJ11806 (FLJ11806), mRNA. /FEA=mRNA /GEN=FLJ11806 /PROD=hypothetical protein FLJ11806 /DB_XREF=gi:13376221 /UG=Hs.323443 hypothetical protein FLJ11806 /FL=gb:NM_024824.1	NM_024824		NP_079100

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204218_at	0.001148	gb:NM_014042.1 /DEF=Homo sapiens DKFZP564M082 protein (DKFZP564M082), mRNA. /FEA=mRNA /GEN=DKFZP564M082 /PROD=DKFZP564M082 protein /DB_XREF=gi:7661621 /UG=Hs.38044 DKFZP564M082 protein /FL=gb:BC005156.1 gb:BC005393.1 gb:AF077206.1 gb:AL080071.1 gb:NM_014042.1	NM_014042		NP_054761
204219_s_at	0.001763	gb:NM_002802.1 /DEF=Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 1 (PSMC1), mRNA. /FEA=mRNA /GEN=PSMC1 /PROD=proteasome (prosome, macropain) 26S subunit,ATPase, 1 /DB_XREF=gi:4506206 /UG=Hs.4745 proteasome (prosome, macropain) 26S subunit, ATPase, 1 /FL=gb:L02426.1 gb:NM_002802.1	NM_002802		NP_002793
204220_at	0.032771	gb:NM_004877.1 /DEF=Homo sapiens glia maturation factor, gamma (GMFG), mRNA. /FEA=mRNA /GEN=GMFG /PROD=glia maturation factor, gamma /DB_XREF=gi:4758439 /UG=Hs.5210 glia maturation factor, gamma /FL=gb:AB001993.1 gb:AF038956.1 gb:NM_004877.1	NM_004877		NP_004868

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
204225_at	7.25E-04	gb:NM_006037.2 /DEF=Homo sapiens histone deacetylase 4 (HDAC4), mRNA. /FEA=mRNA /GEN=HDAC4 /PROD=histone deacetylase 4 /DB_XREF=gi:13259519 /UG=Hs.91400 histone deacetylase 4 /FL=gb:NM_006037.2 gb:AF132607.1	NM_006037		NP_006028
204226_at	0.010065	gb:NM_014393.1 /DEF=Homo sapiens staufer (Drosophila, RNA-binding protein) homolog 2 (STAU2), mRNA. /FEA=mRNA /GEN=STAU2 /PROD=staufer homolog 2 /DB_XREF=gi:7657624 /UG=Hs.96870 staufer (Drosophila, RNA-binding protein) homolog 2 /FL=gb:NM_014393.1	NM_014393		NP_055208
204227_s_at	0.009329	gb:NM_004614.1 /DEF=Homo sapiens thymidine kinase 2, mitochondrial (TK2), mRNA. /FEA=mRNA /GEN=TK2 /PROD=thymidine kinase 2, mitochondrial /DB_XREF=gi:10281329 /UG=Hs.274701 thymidine kinase 2, mitochondrial /FL=gb:NM_004614.1 gb:U77088.1	NM_004614		NP_004605
204232_at	7.63E-04	gb:NM_004106.1 /DEF=Homo sapiens Fc fragment of IgE, high affinity I, receptor for, gamma polypeptide (FCER1G), mRNA. /FEA=mRNA /GEN=FCER1G /PROD=Fc fragment of IgE, high affinity I, receptor for, gamma polypeptide precursor /DB_XREF=gi:4758343 /UG=Hs.743 Fc fragment of IgE, high affinity I, receptor for, gamma polypeptide /FL=gb:M33195.1 gb:NM_004106.1	NM_004106		NP_004097

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204236_at	7.63E-04	gb:NM_002017.2 /DEF=Homo sapiens Friend leukemia virus integration 1 (FLI1), mRNA. /FEA=mRNA /GEN=FLI1 /PROD=Friend leukemia virus integration 1. /DB_XREF=gi:7110592 /UG=Hs.108043 Friend leukemia virus integration 1 /FL=gb:BC001670.1 gb:M98833.3 gb:NM_002017.2	NM_002017		NP_002008
204244_s_at	0.007591	gb:NM_006716.1 /DEF=Homo sapiens activator of S phase kinase (ASK), mRNA. /FEA=mRNA /GEN=ASK /PROD=activator of S phase kinase /DB_XREF=gi:5729733 /UG=Hs.152759 activator of S phase kinase. /FL=gb:AB028069.1 gb:AF160249.1 gb:AF160876.1 gb:NM_006716.1	NM_006716		NP_006707
204246_s_at	0.006722	gb:NM_007234.2 /DEF=Homo sapiens dynactin 3 (p22) (DCTN3), transcript variant 1, mRNA. /FEA=mRNA /GEN=DCTN3 /PROD=dynactin 3, isoform 1. /DB_XREF=gi:13259516 /UG=Hs.15961 dynactin 3 (p22) /FL=gb:NM_007234.2 gb:AF082513.1	NM_007234		NP_077324
204249_s_at	0.004519	gb:NM_005574.2 /DEF=Homo sapiens LIM domain only 2 (rhombotin-like 1) (LMO2), mRNA. /FEA=mRNA /GEN=LMO2 /PROD=LIM domain only 2 /DB_XREF=gi:6633806 /UG=Hs.184585 LIM domain only 2 (rhombotin-like 1) /FL=gb:NM_005574.2	NM_005574		NP_005565

G n s Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204254_s_at	7.63E-04	gb:NM_000376.1 /DEF=Homo sapiens vitamin D (1,25-dihydroxyvitamin D3) receptor (VDR), mRNA. /FEA=mRNA /GEN=VDR /PROD=vitamin D (1,25-dihydroxyvitamin D3) receptor /DB_XREF=gi:4507882 /UG=Hs.2062 vitamin D (1,25-dihydroxyvitamin D3) receptor /FL=gb:AF026260.1 gb:J03258.1 gb:NM_000376.1	NM_000376		NP_000367
204257_at	0.016468	gb:NM_021727.1 /DEF=Homo sapiens fatty acid desaturase 3 (FADS3), mRNA. /FEA=mRNA /GEN=FADS3 /PROD=fatty acid desaturase 3 /DB_XREF=gi:13375615 /UG=Hs.21765 fatty acid desaturase 3 /FL=gb:AF084560.1 gb:NM_021727.1 gb:BC004901.1 gb:AF134404.1	NM_021727		NP_068373
204264_at	0.006031	gb:NM_000098.1 /DEF=Homo sapiens carnitine palmitoyltransferase II (CPT2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=CPT2 /PROD=carnitine palmitoyltransferase II /DB_XREF=gi:4503022 /UG=Hs.274336 carnitine palmitoyltransferase II /FL=gb:U09648.1 gb:BC002445.1 gb:BC005172.1 gb:M58581.1 gb:NM_000098.1	NM_000098		NP_000089
204269_at	0.039362	gb:NM_006875.1 /DEF=Homo sapiens pim-2 oncogene (PIM2), mRNA. /FEA=mRNA /GEN=PIM2 /PROD=pim-2 oncogene /DB_XREF=gi:5803124 /UG=Hs.80205 pim-2 oncogene /FL=gb:U77735.1 gb:NM_006875.1	NM_006875		NP_006866

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204275_at	0.007803	small optic lobes homolog (Drosophila)	AI796687	Hs.55836	NP_005623
204287_at	0.034317	gb:NM_004711.1 /DEF=Homo sapiens synaptogyrin 1 (SYNGR1), mRNA. /FEA=mRNA /GEN=SYNGR1 /PROD=synaptogyrin 1 /DB_XREF=gi:4759199 /UG=Hs.6139 synaptogyrin 1 /FL=gb:NM_004711.1	NM_004711		NP_663791
204295_at	0.024689	gb:NM_003172.1 /DEF=Homo sapiens surfait 1 (SURF1), mRNA. /FEA=mRNA /GEN=SURF1 /PROD=surfait 1 /DB_XREF=gi:4507318 /UG=Hs.3196 surfait 1 /FL=gb:NM_003172.1	NM_003172		NP_003163
204297_at	0.001719	gb:NM_002647.1 /DEF=Homo sapiens phosphoinositide-3-kinase, class 3 (PIK3C3), mRNA. /FEA=mRNA /GEN=PIK3C3 /PROD=phosphoinositide-3-kinase, class 3 /DB_XREF=gi:4505800 /UG=Hs.32971 phosphoinositide-3-kinase, class 3 /FL=gb:NM_002647.1	NM_002647		NP_002638
204307_at	0.014096	Consensus includes gb:AB002295.1 /DEF=Human mRNA for KIAA0297 gene, partial cds. /FEA=mRNA /GEN=KIAA0297 /DB_XREF=gi:2224534 /UG=Hs.11711 KIAA0329 gene product /FL=gb:AB002327.1 gb:NM_014844.1	AB002295		NP_055659

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204308_s_at	0.002603	gb:NM_014844.1 /DEF=Homo sapiens KIAA0329 gene product (KIAA0329), mRNA. /FEA=mRNA /GEN=KIAA0329 /PROD=KIAA0329 gene product /DB_XREF=gi:7662057 /UG=Hs.11711 KIAA0329 gene product /FL=gb:AB002327.1 gb:NM_014844.1	NM_014844		NP_055659
204313_s_at	0.021852	Consensus includes gb:AA161486 /FEA=EST /DB_XREF=gi:1735796 /DB_XREF=est:zq42d09.s1 /CLONE=IMAGE:632369 /UG=Hs.79194 cAMP responsive element binding protein 1 /FL=gb:M27691.1 gb:NM_004379.1	NM_004379		NP_604391
204314_s_at	0.003373	gb:NM_004379.1 /DEF=Homo sapiens cAMP responsive element binding protein 1 (CREB1), mRNA. /FEA=mRNA /GEN=CREB1 /PROD=cAMP responsive element binding protein 1 /DB_XREF=gi:4758053 /UG=Hs.79194 cAMP responsive element binding protein 1 /FL=gb:M27691.1 gb:NM_004379.1	NM_004379		NP_604391
204316_at	0.042858	Consensus includes gb:W19676 /FEA=EST /DB_XREF=gi:1295644 /DB_XREF=est:zb36h07.r1 /CLONE=IMAGE:305725 /UG=Hs.82280 regulator of G-protein signalling 10 /FL=gb:NM_002925.2 gb:AF045229.1	NM_002925		NP_002916

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204336_s_at	0.008944	gb:NM_005873.1 /DEF=Homo sapiens regulator of G-protein signalling 19 (RGS19), mRNA. /FEA=mRNA /GEN=RGS19 /PROD=G protein signalling regulator 19 /DB_XREF=gi:5031704 /UG=Hs.22698 regulator of G-protein signalling 19 /FL=gb:NM_005873.1	NM_005873		NP_005864
204342_at	0.001852	gb:NM_013386.1 /DEF=Homo sapiens hypothetical protein (DKFZp586G0123), mRNA. /FEA=mRNA /GEN=DKFZp586G0123 /PROD=hypothetical protein /DB_XREF=gi:9558726 /UG=Hs.24713 hypothetical protein /FL=gb:AL050209.1 gb:NM_013386.1	NM_013386		NP_037518
204351_at	0.024696	gb:NM_005980.1 /DEF=Homo sapiens S100 calcium-binding protein P (S100P), mRNA. /FEA=mRNA /GEN=S100P /PROD=S100 calcium-binding protein P /DB_XREF=gi:5174662 /UG=Hs.2962 S100 calcium-binding protein P /FL=gb:NM_005980.1	NM_005980		NP_005971
204361_s_at	0.010277	gb:AB014486.1 /DEF=Homo sapiens mRNA for RA70, complete cds. /FEA=mRNA /GEN=RA70 /PROD=RA70 /DB_XREF=gi:4062959 /UG=Hs.52644 SKAP55 homologue /FL=gb:BC002893.1 gb:AF072166.1 gb:AB014486.1 gb:AF051323.1 gb:NM_003930.1	AB014486		NP_003921

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unique Accession No.	Protein Accession No.
204362_at	0.00109	gb:NM_003930.1 /DEF=Homo sapiens SKAP55 homologue (SKAP-HOM), mRNA. /FEA=mRNA /GEN=SKAP-HOM /PROD=SKAP55 homologue /DB_XREF=gi:4506962 /UG=Hs.52644 SKAP55 homologue /FL=gb:BC002893.1 gb:AF072166.1 gb:AB014486.1 gb:AF051323.1 gb:NM_003930.1	NM_003930		NP_003921
204366_s_at	0.004403	gb:NM_001521.1 /DEF=Homo sapiens general transcription factor IIIC, polypeptide 2 (beta subunit, 110kD), (GTF3C2), mRNA. /FEA=mRNA /GEN=GTF3C2 /PROD=general transcription factor IIIC, polypeptide 2(beta subunit, 110kD) /DB_XREF=gi:4504204 /UG=Hs.75782 general transcription factor IIIC, polypeptide 2 (beta subunit, 110kD) /FL=gb:D13636.1 gb:NM_001521.1	NM_001521		NP_001512
204370_at	7.25E-04	gb:NM_006831.1 /DEF=Homo sapiens ATPGTP-binding protein (HEAB), mRNA. /FEA=mRNA /GEN=HEAB /PROD=ATPGTP-binding protein /DB_XREF=gi:5803028 /UG=Hs.87465 ATPGTP-binding protein /FL=gb:BC000446.1 gb:U73524.1 gb:NM_006831.1	NM_006831		NP_006822

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204387_x_at	0.008944	gb:NM_024026.1 /DEF=Homo sapiens hypothetical protein MGC3243 (MGC3243), mRNA. /FEA=mRNA /GEN=MGC3243 /PROD=hypothetical protein MGC3243 /DB_XREF=gi:13128969 /UG=Hs.182695 hypothetical protein MGC3243 /FL=gb:BC000002.1 gb:NM_024026.1	NM_024026		NP_076931
204391_x_at	0.017687	gb:NM_015905.1 /DEF=Homo sapiens transcriptional intermediary factor 1 (TIF1), mRNA. /FEA=mRNA /GEN=TIF1 /PROD=transcriptional intermediary factor 1 alpha /DB_XREF=gi:7706233 /UG=Hs.183858 transcriptional intermediary factor 1 /FL=gb:AF009353.1 gb:AF119042.1 gb:NM_003852.1 gb:NM_015905.1	NM_015905		
204393_s_at	0.001432	gb:NM_001099.2 /DEF=Homo sapiens acid phosphatase, prostate (ACPP), mRNA. /FEA=mRNA /GEN=ACPP /PROD=prostatic acid phosphatase precursor /DB_XREF=gi:6382063 /UG=Hs.1852 acid phosphatase, prostate /FL=gb:M24902.1 gb:M34840.1 gb:NM_001099.2	NM_001099		NP_001090

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
204409_s_at	0.007659	gb:BC005248.1 /DEF=Homo sapiens, clone MGC:12282, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:12282) /DB_XREF=gi:13528902 /UG=Hs.155103 eukaryotic translation initiation factor 1A, Y chromosome /FL=gb:BC005248.1 gb:AF000987.1 gb:NM_004681.1	BC005248		NP_004672
204411_at	0.011797	gb:NM_017596.1 /DEF=Homo sapiens KIAA0449 protein (KIAA0449), mRNA. /FEA=mRNA /GEN=KIAA0449 /PROD=hypothetical protein DKFZp434J212 /DB_XREF=gi:8922142 /UG=Hs.169182 KIAA0449 protein /FL=gb:NM_017596.1	NM_017596		NP_060066
204417_at	7.63E-04	gb:NM_000153.1 /DEF=Homo sapiens galactosylceramidase (Krabbe disease) (GALC), mRNA. /FEA=mRNA /GEN=GALC /PROD=galactosylceramidase precursor /DB_XREF=gi:4557612 /UG=Hs.273 galactosylceramidase (Krabbe disease) /FL=gb:L23116.1 gb:NM_000153.1 gb:D25283.1	NM_000153		NP_000144
204419_x_at	7.55E-04	gb:NM_000184.1 /DEF=Homo sapiens hemoglobin, gamma G (HBG2), mRNA. /FEA=mRNA /GEN=HBG2 /PROD=hemoglobin, gamma G /DB_XREF=gi:6715606 /UG=Hs.283108 hemoglobin, gamma G /FL=gb:NM_000184.1	NM_000184		NP_000175

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204423_at	0.019229	gb:NM_013255.1 /DEF=Homo sapiens muskelin 1, intracellular mediator containing kelch motifs (MKLN1), mRNA. /FEA=mRNA /GEN=MKLN1 /PROD=muskelin 1, intracellular mediator containing kelch motifs /DB_XREF=gi:7019458 /UG=Hs.288791 muskelin 1, intracellular mediator containing kelch motifs /FL=gb:BC002834.1 gb:AF047489.1 gb:NM_013255.1	NM_013255		NP_037387
204427_s_at	0.047575	gb:NM_006815.1 /DEF=Homo sapiens coated vesicle membrane protein (RNP24), mRNA. /FEA=mRNA /GEN=RNP24 /PROD=coated vesicle membrane protein /DB_XREF=gi:5803148 /UG=Hs.323378 coated vesicle membrane protein. /FL=gb:NM_006815.1	NM_006815		NP_006806
204445_s_at	0.015637	arachidonate 5-lipoxygenase	AI361850	Hs.89499	NP_000689
204446_s_at	0.007174	gb:NM_000698.1 /DEF=Homo sapiens arachidonate 5-lipoxygenase (ALOX5), mRNA. /FEA=mRNA /GEN=ALOX5 /PROD=arachidonate 5-lipoxygenase /DB_XREF=gi:4502056 /UG=Hs.89499 arachidonate 5-lipoxygenase /FL=gb:J03600.1 gb:J03571.1 gb:NM_000698.1	NM_000698		NP_000689
204466_s_at	7.63E-04	synuclein, alpha (non A4 component of amyloid precursor)	BG260394	Hs.76930	NP_009292

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204467_s_at	0.003706	gb:NM_000345.2 /DEF=Homo sapiens synuclein, alpha (non A4 component of amyloid precursor) (SNCA), transcript variant NACP140, mRNA. /FEA=mRNA /GEN=SNCA /PROD=alpha-synuclein isoform NACP140 /DB_XREF=gi:6806896 /UG=Hs.76930 synuclein, alpha (non A4 component of amyloid precursor) /FL=gb:L08850.1 gb:NM_000345.2	NM_000345		NP_009292
204478_s_at	7.41E-04	gb:NM_002871.1 /DEF=Homo sapiens RAB interacting factor (RABIF), mRNA. /FEA=mRNA /GEN=RABIF /PROD=RAB interacting factor /DB_XREF=gi:4506378 /UG=Hs.90875 RAB interacting factor /FL=gb:U74324.1 gb:NM_002871.1	NM_002871		NP_002862
204489_s_at	0.006483	gb:NM_000610.1 /DEF=Homo sapiens CD44 antigen (homing function and Indian blood group system) (CD44), mRNA. /FEA=mRNA /GEN=CD44 /PROD=CD44 antigen (homing function and Indian bloodgroup system) /DB_XREF=gi:10835162 /UG=Hs.169610 CD44 antigen (homing function and Indian blood group system) /FL=gb:NM_000610.1 gb:U40373.1 gb:M59040.1 gb:M24915.1	NM_000610		NP_000601
204491_at	0.047555	phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 duncce homolog, Drosophila)	R40917	Hs.172081	NP_006194

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladd rCancer					
Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204496_at	0.008874	gb:NM_014574.1 /DEF=Homo sapiens nuclear autoantigen (GS2NA), mRNA. /FEA=mRNA /GEN=GS2NA /PROD=nuclear autoantigen /DB_XREF=gi:11128016 /UG=Hs.183105 nuclear autoantigen /FL=gb:NM_014574.1 gb:U17989.1	NM_014574		NP_055389
204500_s_at	7.63E-04	gb:NM_015239.1 /DEF=Homo sapiens KIAA1035 protein (KIAA1035), mRNA. /FEA=mRNA /GEN=KIAA1035 /PROD=KIAA1035 protein /DB_XREF=gi:7662451 /UG=Hs.21542 KIAA1035 protein /FL=gb:NM_015239.1	NM_015239		NP_056054
204505_s_at	7.63E-04	gb:NM_001978.1 /DEF=Homo sapiens erythrocyte membrane protein band 4.9 (dematin) (EPB49), mRNA. /FEA=mRNA /GEN=EPB49 /PROD=erythrocyte membrane protein band 4.9 (dematin) /DB_XREF=gi:4503580 /UG=Hs.274122 erythrocyte membrane protein band 4.9 (dematin) /FL=gb:NM_001978.1 gb:U28389.1	NM_001978		NP_001969

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204507_s_at	0.013527	gb:NM_000945.1 /DEF=Homo sapiens protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I) (PPP3R1), mRNA. /FEA=mRNA /GEN=PPP3R1 /PROD=protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I) /DB_XREF=gi:4506024 /UG=Hs.278540 protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I) /FL=gb:M30773.1 gb:NM_000945.1	NM_000945		NP_000936
204513_s_at	7.25E-04	gb:NM_014800.1 /DEF=Homo sapiens KIAA0281 gene product (KIAA0281), mRNA. /FEA=mRNA /GEN=KIAA0281 /PROD=KIAA0281 gene product /DB_XREF=gi:7662037 /UG=Hs.31463 KIAA0281 gene product /FL=gb:BC003051.1 gb:D87457.1 gb:NM_014800.1	NM_014800		NP_569709
204526_s_at	7.63E-04	gb:NM_007063.1 /DEF=Homo sapiens vascular Rab-GAPTBC-containing (VRP), mRNA. /FEA=mRNA /GEN=VRP /PROD=vascular Rab-GAPTBC-containing /DB_XREF=gi:5902153 /UG=Hs.164170 vascular Rab-GAPTBC-containing /FL=gb:AB024057.1 gb:NM_007063.1	NM_007063		NP_008994

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gen Identifier	p-value	Description	Gen Accession No.	Unig ne Accession No.	Protein Accession No.
204527_at	0.011487	gb:NM_000259.1 /DEF=Homo sapiens myosin VA (heavy polypeptide 12, myoxin) (MYO5A), mRNA. /FEA=mRNA /GEN=MYO5A /PROD=myosin VA (heavy polypeptide 12, myoxin) /DB_XREF=gi:10835118 /UG=Hs.170157 myosin VA (heavy polypeptide 12, myoxin) /FL=gb:NM_000259.1 gb:U90942.1	NM_000259		NP_000250
204546_at	6.66E-04	gb:NM_014732.1 /DEF=Homo sapiens KIAA0513 gene product (KIAA0513), mRNA. /FEA=mRNA /GEN=KIAA0513 /PROD=KIAA0513 gene product /DB_XREF=gi:7662163 /UG=Hs.301658 KIAA0513 gene product /FL=gb:AB011085.1 gb:NM_014732.1	NM_014732		NP_055547
204554_at	7.41E-04	Consensus includes gb:AL109928 /DEF=Human DNA sequence from clone RP4-551D2 on chromosome 20q13.2-13.33: Contains the gene for a novel Cadherin domain protein, a novel gene, the PPP1R6 gene for protein phosphatase 1 regulatory subunit 6, the 5 end of the SYCP2 gene for synaptonemal... /FEA=mRNA_6 /DB_XREF=gi:7981303 /UG=Hs.42215 protein phosphatase 1, regulatory subunit 6 /FL=gb:NM_006242.2	AL109928		

Genes Corresponding To Differentially Expressed Genes in Figure 18 - BladderCancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204565_at	7.41E-04	gb:NM_018473.1 /DEF=Homo sapiens uncharacterized hypothalamus protein HT012 (HT012), mRNA. /FEA=mRNA /GEN=HT012 /PROD=uncharacterized hypothalamus protein HT012 /DB_XREF=gi:8923811 /UG=Hs.9676 uncharacterized hypothalamus protein HT012 /FL=gb:BC000894.1 gb:AF274952.1 gb:AF220186.1 gb:NM_018473.1	NM_018473		NP_060943
204599_s_at	0.006622	gb:NM_006428.1 /DEF=Homo sapiens melanoma-associated antigen recognised by cytotoxic T lymphocytes (MAAT1), mRNA. /FEA=mRNA /GEN=MAAT1 /PROD=melanoma-associated antigen recognised by T lymphocytes /DB_XREF=gi:5453835 /UG=Hs.279869 melanoma-associated antigen recognised by cytotoxic T lymphocytes /FL=gb:BC000507.1 gb:BC000990.2 gb:NM_006428.1 gb:U19796.1	NM_006428		NP_006419
204604_at	0.00121	gb:NM_012395.1 /DEF=Homo sapiens PFTAIRE protein kinase 1 (PFTK1), mRNA. /FEA=mRNA /GEN=PFTK1 /PROD=PFTAIRE protein kinase 1 /DB_XREF=gi:6912583 /UG=Hs.57856 PFTAIRE protein kinase 1 /FL=gb:AB020641.1 gb:NM_012395.1	NM_012395		NP_036527

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204613_at	0.001942	gb:NM_002661.1 /DEF=Homo sapiens phospholipase C, gamma 2 (phosphatidylinositol-specific) (PLCG2), mRNA. /FEA=mRNA /GEN=PLCG2 /PROD=phospholipase C, gamma 2(phosphatidylinositol-specific) /DB_XREF=gi:4505870 /UG=Hs.75648 phospholipase C, gamma 2 (phosphatidylinositol-specific) /FL=gb:M37238.1 gb:NM_002661.1	NM_002661		NP_002652
204614_at	7.34E-04	gb:NM_002575.1 /DEF=Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2 (SERPINB2), mRNA. /FEA=mRNA /GEN=SERPINB2 /PROD=serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2 /DB_XREF=gi:4505594 /UG=Hs.75716 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2 /FL=gb:J02685.1 gb:J03603.1 gb:M18082.1 gb:NM_002575.1	NM_002575		NP_002566
204615_x_at	6.66E-04	gb:NM_004508.1 /DEF=Homo sapiens isopentenyl-diphosphate delta isomerase (IDI1), mRNA. /FEA=mRNA /GEN=IDI1 /PROD=isopentenyl-diphosphate delta isomerase /DB_XREF=gi:4758583 /UG=Hs.76038 isopentenyl-diphosphate delta isomerase /FL=gb:NM_004508.1	NM_004508		NP_004499

Genes Corresponding To Differentially Expressed Genes in Figure 18 - Bladder Cancer					
Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204616_at	0.017445	gb:NM_006002.1 /DEF=Homo sapiens ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase) (UCHL3), mRNA. /FEA=mRNA /GEN=UCHL3 /PROD=ubiquitin carboxyl-terminal esterase L3(ubiquitin thiolesterase) /DB_XREF=gi:5174740 /UG=Hs.77917 ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase) /FL=gb:M30496.1 gb:NM_006002.1	NM_006002		NP_005993
204619_s_at	0.007174	chondroitin sulfate proteoglycan 2 (versican)	BF590263	Hs.81800	NP_004376
204620_s_at	7.63E-04	gb:NM_004385.1 /DEF=Homo sapiens chondroitin sulfate proteoglycan 2 (versican) (CSPG2), mRNA. /FEA=mRNA /GEN=CSPG2 /PROD=chondroitin sulfate proteoglycan 2 (versican) /DB_XREF=gi:4758081 /UG=Hs.81800 chondroitin sulfate proteoglycan 2 (versican) /FL=gb:NM_004385.1	NM_004385		NP_004376
204630_s_at	0.017845	gb:NM_004871.1 /DEF=Homo sapiens golgi SNAP receptor complex member 1 (GOSR1), mRNA. /FEA=mRNA /GEN=GOSR1 /PROD=golgi SNAP receptor complex member 1 /DB_XREF=gi:4758455 /UG=Hs.8868 golgi SNAP receptor complex member 1 /FL=gb:AF073926.1 gb:NM_004871.1	NM_004871		NP_004862